

Spatiotemporal motion estimation for respiratory-correlated imaging of the lungs

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Purpose: Four-dimensional computed tomography (4D CT) can provide patient-specific motion information for radiotherapy planning and delivery. Motion estimation in 4D CT is challenging due to the reduced image quality and the presence of artifacts. We aim to improve the robustness of deformable registration applied to respiratory-correlated imaging of the lungs, by using a global problem formulation and pursuing a restrictive parametrization for the spatiotemporal deformation model.

Methods: A spatial transformation based on free-form deformations was extended to the temporal domain, by explicitly modeling the trajectory using a cyclic temporal model based on B-splines. A global registration criterion allowed to consider the entire image sequence simultaneously and enforce the temporal coherence of the deformation throughout the respiratory cycle. To ensure a parametrization capable of capturing the dynamics of respiratory motion, a prestudy was performed on the temporal dimension separately. The temporal parameters were tuned by fitting them to diaphragm motion data acquired for a large patient group. Suitable properties were retained and applied to spatiotemporal registration of 4D CT data. Registration results were validated using large sets of landmarks and compared to consecutive spatial registrations. To illustrate the benefit of the spatiotemporal approach, we also assessed the performance in the presence of motion-induced artifacts.

Results: Cubic B-splines gave better or similar fitting results as lower orders and were selected because of their inherently stronger regularization. The fitting and registration errors increased gradually with the temporal control point spacing, representing a trade-off between achievable accuracy and sensitivity to noise and artifacts. A piecewise smooth trajectory model, allowing for a discontinuous change of speed at end-inhale, was found most suitable to account for the sudden changes of motion at this breathing phase. The spatiotemporal modeling allowed a reduction of the number of parameters of 45%, while maintaining registration accuracy within 0.1 mm. The approach reduced the sensitivity to artifacts.

Conclusions: Spatiotemporal registration can provide accurate motion estimation for 4D CT and improves the robustness to artifacts. © 2011 American Association of Physicists in Medicine. [DOI: [10.1118/1.3523619](https://doi.org/10.1118/1.3523619)]

Key words: deformable registration, respiratory motion, 4D CT

I. INTRODUCTION

The advent of four-dimensional (4D) computed tomography (CT) has allowed patient-specific respiratory motion infor-

mation to be incorporated into radiation therapy planning and delivery. 4D CT provides multiple three-dimensional (3D) CT volumes, representing the patient at different stages

of the breathing cycle.¹⁻⁴ The additional patient data imply an order of magnitude increase in the workload required to obtain a 4D treatment plan. Deformable registration is the tool that can facilitate partial automation of the 4D planning process.⁵ It can provide the motion fields which are required for automating tasks such as recontouring of anatomic structures,⁶ patient-specific margin definition,⁷ or 4D treatment plan evaluation.⁸ Deformable image registration is also an enabling tool for alternative applications of respiratory-correlated imaging such as ventilation imaging,⁹ motion compensation,^{10,11} or motion modeling.^{12,13} Although extensive validation is required before extending the clinical use of deformable image registration, it is expected to become a standard methodology in radiotherapy.^{14,15}

Deformable image registration can be described as the task of finding a suitable geometric transformation between corresponding image data, such that a transformed image becomes similar to another one.¹⁶ While the concept of image registration is easily described, the underlying numerical problem is difficult to solve. Mainly because the registration problem is *ill-posed*. Small changes of the input images may lead to very different registration results. Moreover, the solution might not be unique. Salient image information might be sparse or ambiguous, and the acquisition process might have introduced noise and artifacts. To facilitate the process, prior knowledge about the deformation should be incorporated in the registration framework in order to favor solutions with plausible physical characteristics. Explicit parametric restrictions can constrain the optimization to transformations that represent suitable properties. This approach can offer a reduction of the search space by making the description more problem-specific and consequently improve the robustness of the optimization process.

An example are spatiotemporal registration schemes, which consist in a global formulation of the motion estimation problem for temporal image sequences. Rather than estimating frame-to-frame displacements individually, the entire sequence is considered simultaneously, allowing to enforce the temporal coherence of the deformation across the sequence. By making assumptions such as smoothness about the temporal variations of the transformation, these approaches often enable a more compact and restrictive description of the full motion estimation problem. Spatiotemporal deformable registration has received considerable attention in literature, mostly in cardiac image analysis,¹⁷⁻²⁶ but more recently also for respiratory-correlated imaging of the thorax.²⁷⁻²⁹ Usually, a 3D-4D formulation is utilized to find a smooth time-dependent deformation field that aligns all images from a given input sequence with a reference image, which can be a frame of that same sequence.^{18,22-24,27,29} Sometimes, spatial as well as temporal alignment of multiple image sequences is desirable, leading to a 4D-4D registration framework.^{20,21,25,28}

In comparison to conventional diagnostic CT, 4D CT images tend to be acquired at lower spatial resolution and are characterized by higher noise levels because of the low radiation dose per image. In addition, an alarmingly high number of acquisitions contains motion-induced artifacts,³⁰

mainly due to irregular patient breathing during image acquisition. In the case of artifacts, the image information can be considered locally invalid, as it does not correspond to the patient anatomy. Clinical use of the estimated motion fields requires them to be as close to the unknown reality as possible. A problem-specific, spatiotemporal deformation model could contribute in reducing sensitivity to local image irregularities and render the motion estimate more plausible and potentially more representative of the patient's breathing motion under these challenging circumstances.

In this study, we develop a spatiotemporal registration scheme for lung motion quantification in respiratory-correlated sequences. Our primary objective is to obtain a low-dimensional representation of the 4D deformation model, capable of accurately representing the respiratory motion, while being more robust to artifacts and increased noise levels. The approach consists of a 3D-4D problem formulation in which temporal regularization is pursued by explicitly modeling the trajectory of moving structures. With respect to previous work on spatiotemporal registration, we specifically focus on respiratory-correlated image sequences and develop and evaluate a cyclic trajectory model for representing the motion over an entire breathing cycle. In addition, the chosen parametrization reflects our aim to improve registration robustness by rendering the deformation model more problem-specific.

II. METHOD

The spatiotemporal transformation will be developed incrementally. We first describe a conventional spatial registration, of which the proposed method can be seen as an extension. Next, the temporal dimension is considered separately and the method for modeling the trajectory is detailed. The sought spatiotemporal deformation function is obtained by combining both.

II.A. Problem description

Consider a 4D sequence, represented by an intensity function $f(\mathbf{i}, k) \in \mathbb{R}$ with $\mathbf{i} \in \mathbf{I} \subset \mathbb{Z}^3$ and $k \in \mathbf{K} \subset \mathbb{Z}$; \mathbf{I} and \mathbf{K} being the set of spatial and temporal sample indices, respectively. We wish to analyze the motion with respect to the 3D reference frame at time index $k_r \in \mathbf{K}$. The task of motion estimation throughout the 4D sequence is formulated as the search for the unknown spatiotemporal transformation \mathcal{T}_{st} , defined for $\mathbf{I} \times \mathbf{K} \mapsto \mathbb{R}^3$, where $\mathcal{T}_{st}(\mathbf{i}, k)$ represents the location of a point at time k which was at position \mathbf{i} at time k_r .

II.B. Spatial registration

Consider the subproblem of retrieving the transformation \mathcal{T}_s (in which s stands for spatial) between the reference volume and the frame at time k . A continuous representation is employed for the spatial transformation using free-form deformations based on B-splines,³¹

$$\mathcal{T}_s(\mathbf{x}) = \mathbf{x} + \sum_{j \in \mathbf{J}} \mathbf{a}_j \phi_j(\mathbf{x}), \quad (1)$$

where $\mathbf{x} \in \mathbf{X}$ is the continuous spatial coordinate associated with \mathbf{I} , $\mathbf{J} \subset \mathbb{Z}^3$ is the set of spatial parameter indices considered for basis functions $\phi_j(\mathbf{x}) = \beta^n(\mathbf{x}/h - \mathbf{j})$ with $h \in \mathbb{R}$ the uniform spatial control point spacing, and β^n the tensor product of one-dimensional B-spline kernels of degree n . We used cubic B-splines for the spatial basis functions ($n=3$). The parameters of \mathcal{T}_s are the B-spline coefficients $\mathbf{a}_j \in \mathbb{R}^3$ (one for each component of the deformation), i.e., \mathcal{T}_s is fully characterized by specifying $\mathbf{a} = \{\mathbf{a}_j\}_{j \in \mathbf{J}}$.

We define a similarity criterion \mathcal{J}_s , based on the mean squared intensity differences with respect to the samples of the reference volume,

$$\mathcal{J}_s(\mathcal{T}_s, k) = \frac{1}{N_{\mathbf{I} \in \mathbf{I}}} \sum (f(\mathcal{T}_s(i), k) - f(i, k_r))^2, \quad (2)$$

with $N_{\mathbf{I}}$ the number of spatial samples considered. We chose this criterion because of its fast computation time and the smoothness of the resulting search space. For simplicity, no explicit regularization term was included in the criterion. For now, only the influence of the parametrization of the deformation function was explored. Evaluating the intensity function f at nongrid positions requires a continuous representation for which we used cubic B-spline interpolation,

$$f(\mathbf{x}, k) = \sum_{i \in \mathbf{I}} \mathbf{d}_i \beta^n(\mathbf{x} - \mathbf{i}). \quad (3)$$

Coefficients \mathbf{d}_i are found quickly from the image intensities using recursive filtering.³²

Solving the spatial registration problem for frame k comes down to estimating the optimal parameters \mathbf{a}^* in the sense of the criterion \mathcal{J}_s ,

$$\mathbf{a}^* = \arg \min_{\mathbf{a}} \mathcal{J}_s(\mathcal{T}_s; k). \quad (4)$$

By solving Eq. (4) consecutively for all $k \in \mathbf{K}$ except k_r , a solution to the 4D motion estimation problem can be composed. Solutions obtained for previous k values can be used to initialize subsequent registrations.

II.C. Trajectory modeling

Temporal sequences enable modeling the temporal variations of the estimated deformations. Tissue trajectories are expected to evolve smoothly and continuously over time, allowing to introduce constraints which enforce the temporal coherence of the deformation across the sequence. This is similar to the approach described for the spatial dimensions. Nonetheless, the temporal dimension is handled separately as it is inherently different. For instance, in the case of respiratory-correlated CT, the sequence is periodic and the number of temporal samples is low compared to spatial samples.

Trajectory model. Let $t \in \mathbf{T}$ be the continuous coordinate associated with \mathbf{K} and suppose for simplicity $\mathbf{T} = [0, t_e]$. Let $\mathcal{T}_t(\mathbf{x}, t)$ denote the trajectory of a point at position \mathbf{x} at the

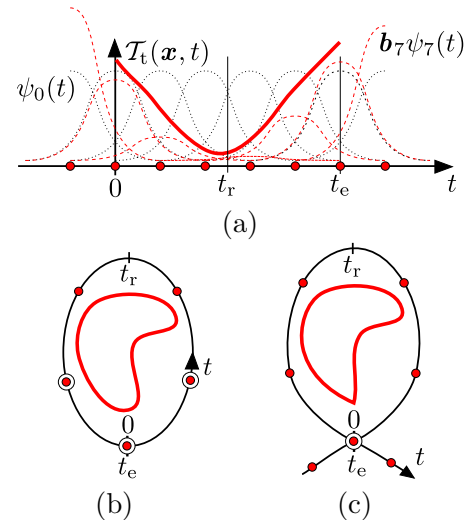


FIG. 1. (a) Schematic representation of a trajectory model based on cubic B-splines, with eight control points (dots) placed uniformly inside and just outside $[0, t_e]$. Each corresponds to a B-spline kernel ψ_l (dotted line) and $\mathcal{T}_t(\mathbf{x}, t)$ (solid curve) is found by combining the scaled kernels (dashed line). [(b) and (c)] Alternative, representation with a cyclic temporal axis wrapped around the trajectory. Large control points indicate a constraint is applied. (b) The smooth trajectory model \mathcal{T}_t and (c) the piecewise smooth trajectory model \mathcal{T}_t^* .

reference time t_r . The search for \mathcal{T}_t is limited to continuous and smooth functions of t , by expressing it using a suitable set of basis functions $\{\psi_l\}_{l \in \mathbf{L}}$,

$$\mathcal{T}_t(\mathbf{x}, t) = \mathbf{x} + \sum_{l \in \mathbf{L}} \mathbf{b}_l \psi_l(t). \quad (5)$$

$\mathbf{L} \subset \mathbb{Z}$ is the set of temporal parameter indices and $\mathbf{b}_l \in \mathbb{R}^3$ the coefficients of the basis functions. We adopted uniform B-spline basis functions^{17,20,22,27} of order $m \in \mathbb{N}$, $\psi_l(t) = \beta^m(t/s - l)$ with $s \in \mathbb{R}$ the temporal control point spacing, because of their good approximation properties, computational simplicity, and implicit smoothness. In Ledesma-Carbayo *et al.*,²² temporal B-splines were found to work at least as well as harmonic functions.^{18,19,23} Figure 1(a) shows a schematic, one-dimensional representation of a trajectory model based on cubic B-splines ($m=3$), with five internal control points ($s=t_e/5$) placed uniformly along the considered interval $[0, t_e]$. Evaluating \mathcal{T}_t near the borders of the interval requires taking into account control points with non-zero weight just outside the interval. It can be seen that a total of eight degrees of freedom is considered, represented by the B-spline coefficients \mathbf{b}_0 to \mathbf{b}_7 .

Smooth trajectory model. The trajectory model can be further constrained by incorporating *a priori* knowledge of the motion, leading to a more restrictive parametrization. For instance, 4D CT data are inherently periodic. In addition, trajectories can be expected to be smooth functions of time. The trajectory can be made periodic and smooth throughout the entire cycle by imposing the same order of smoothness to the endpoints as the rest of the trajectory, thus obtaining $\mathcal{T}_t \in C^{m-1}(\mathbf{T})$. This leads to the set of m conditions

$$\frac{\partial^z T_t(\mathbf{x}, 0)}{\partial t^z} = \frac{\partial^z T_t(\mathbf{x}, t_e)}{\partial t^z} \quad \text{for } z = [0, \dots, m-1]. \quad (6)$$

As will be shown, each condition results in a linear equation for the model parameters, allowing to express one of the parameters in function of the others. A schematic representation of a trajectory satisfying Eq. (6) is shown in Fig. 1(b). A cyclic time axis is shown to illustrate the placement and the influence of the control points. The banded control points indicate that a constraint is applied.

Piecewise smooth trajectory model. Due to the limited temporal resolution of 4D CT, and depending on the degrees of freedom considered in Eq. (5), the smoothness constraint might be too restrictive, leading to locally reduced representation accuracy in regions where the velocity is varying rapidly. This can be the case for the end-inhale phase where fast inversion of the motion takes place. Alternatively, we can locally relax the smoothness constraints and propose a piecewise smooth trajectory representation T_t^* . A similar expansion as Eq. (5) is utilized for T_t^* but, assuming end-inhale corresponds to $t=0$, a single constraint is applied at end-inhale,

$$T_t^*(\mathbf{x}, 0) = T_t^*(\mathbf{x}, t_e). \quad (7)$$

This condition leads to periodic trajectories (C^0 continuity), but allows a discontinuity in the velocity at end-inhale [Fig. 1(c)]. In this case, the sections near end-inhale are parametrized independently which implies a local increase of control points and degrees of freedom.

Temporal constraints. It is instructive to consider the effect of the temporal constraints on the trajectory model. For instance, as deformation is estimated with respect to a reference, by definition

$$T_t(\mathbf{x}, t_r) = \mathbf{x}. \quad (8)$$

This condition allows to express one B-spline coefficient in terms of the others, effectively removing one degree of freedom from the system, i.e.,

$$b_{l_r} = - \sum_{l \in L, l \neq l_r} b_l \frac{\psi_l(t_r)}{\psi_{l_r}(t_r)}, \quad (9)$$

in which we constrained the parameter b_{l_r} associated with the basis function ψ_{l_r} , which is nonzero at t_r . Following Ledesma-Carbayo *et al.*,²² introducing Eq. (9) into Eq. (5) and regrouping terms for each of the parameters allows the temporal model to be expressed using a smaller set of constrained basis functions,

$$\psi_l^c(t) = \psi_l(t) - \frac{\psi_l(t_r)\psi_{l_r}(t)}{\psi_{l_r}(t_r)}, \quad (10)$$

that only generates trajectories that satisfy Eq. (8).

In the following, we will denote L^c the set of temporal indices of basis functions to which constraints (6) and (8) have been applied.

II.D. Spatiotemporal registration

Estimating the motion in a 4D CT sequence by performing consecutive spatial registrations using Eq. (1) fails to exploit the temporal relation between the frames. This is remedied by modeling the trajectory as in Eq. (5). A global approach is found by coupling the temporal and the spatial deformation model,

$$T_{st}(\mathbf{x}, t) = \mathbf{x} + \sum_{j \in J} \sum_{l \in L^c} c_{j,l} \phi_j(\mathbf{x}) \psi_l^c(t). \quad (11)$$

The result is a linear, spatiotemporal deformation function, separable in space and time. A straightforward extension to the temporal dimension of Eq. (2) leads to the criterion

$$\mathcal{J}_{st}(T_{st}) = \frac{1}{N_{\mathbf{K} \times \mathbf{K}}} \sum_{\mathbf{K} \times \mathbf{K}} \mathcal{J}_s(T_{st}, k) \quad (12)$$

to be optimized with respect to the parameters $\mathbf{c} = \{c_{j,l}\}_{j \in J, l \in L^c}$. We will use T_{st} and T_{st}^* in reference to the spatiotemporal deformation models obtained when using the smooth and piecewise smooth temporal models T_t and T_t^* , respectively.

II.E. Motion mask extraction

Breathing motion is characterized by sliding of the liver and lungs, resulting in a discontinuity of the motion at the pleural wall.^{33,34} Accurate matching in these regions requires a complex spatial transformation, even though the remainder of the deformation can be considered smooth. We previously addressed this issue³⁵ by automatically extracting a motion mask, dividing the thorax into moving (lungs, mediastinum, and abdomen), and less-moving regions (the remainder).

Motion masks were computed for all frames of f . The result is the division of the thorax into two subregions $\mathbf{I}_{in}, \mathbf{I}_{out} \subset \mathbf{I}$, roughly representing the inner and outer thoracic structures. For each of the subregions, a separate registration problem can be formulated following the method described by Wu *et al.*,³³ with the advantage that the search can be limited to spatially smooth deformations. In the following, we focused on the inner thoracic structures.

II.F. Optimization

The spatiotemporal approach allows a more restrictive parametrization of the transform and reduces the total number of degrees of freedom of the 4D motion estimation problem compared to consecutively applying T_s . However, directly minimizing Eq. (12) considers all degrees of freedom simultaneously, increasing the dimensionality of the optimization problem with respect to one 3D-3D registration. In response, a multiresolution approach was employed, allowing to gradually increase the complexity of the problem. The resolution of the spatial dimensions of both the image sequence and the B-spline control point grid of the transformation was doubled in each of three consecutive resolution levels. The final image resolution was set to 2 mm. We previously found that, in combination with a motion mask, a control point grid spacing $h=32$ mm provided a good compromise between

registration efficiency and accuracy. The temporal dimension, characterized by low resolution, remained unmodified throughout the optimization.

Each level was handled using a quasi-Newton approach in the form of the limited memory BFGS method³⁶ because of its high precision and improved rate of convergence with respect to simple gradient descent algorithms.³⁷ The procedure started from a zero deformation, and subsequent levels were initialized by upsampling the solution found at the previous level. The required partial derivatives of the similarity criterion can be calculated explicitly, for instance, for \mathcal{T}_{st} ,

$$\frac{\partial \mathcal{J}_{st}(\mathcal{T}_{st})}{\partial \mathbf{c}_{j,l}} = \frac{2}{N_{\mathbf{K}} N_{\mathbf{I}}} \sum_{\mathbf{k} \in \mathbf{K}} \sum_{\mathbf{i} \in \mathbf{I}} (f(\mathcal{T}_{st}(\mathbf{i}, \mathbf{k}), \mathbf{k}) - f(\mathbf{i}, \mathbf{k}_r)) \frac{\partial f(\mathcal{T}_{st}(\mathbf{i}, \mathbf{k}), \mathbf{k})}{\partial \mathbf{x}} \frac{\partial \mathcal{T}_{st}(\mathbf{i}, \mathbf{k})}{\partial \mathbf{c}_{j,l}}. \quad (13)$$

$\partial f / \partial \mathbf{x}$ is found by deriving Eq. (3), while $\partial \mathcal{T}_{st} / \partial \mathbf{c}_{j,l}$ can be calculated considering Eq. (11). For instance, for the q -th spatial component,

$$\frac{\partial f(\mathbf{x}, \mathbf{k})}{\partial x_q} = \sum_{\mathbf{i} \in \mathbf{I}} \mathbf{d}_{\mathbf{i}} \frac{\partial \beta^n(x_q)}{\partial x_q} \prod_{\zeta \neq q} \beta^n(x_{\zeta} - i_{\zeta}), \quad (14)$$

$$\frac{\partial \mathcal{T}_{st}(\mathbf{i}, \mathbf{k})}{\partial \mathbf{c}_{j,l,q}} = \phi_j(\mathbf{x}) \psi_l^q(t). \quad (15)$$

II.G. Implementation

The registration algorithms were implemented in C++. Evaluating $\mathcal{T}_{st}(\mathbf{i}, \mathbf{k})$ was performed using B-LUTs:³⁸ a fast, low memory B-spline implementation based on a look-up table of B-spline tensor products $\beta^n(\mathbf{x})$. Registration algorithms were multithreaded and executed on an eight-core system. The execution times depended on the specifics of the 4D CT data set. Consecutively registering all frames of f using \mathcal{T}_s required between 5 and 10 hours, whereas \mathcal{T}_{st} and \mathcal{T}_{st}^* required about twice as much time.

In comparison, the most expensive step for the spatiotemporal approach is the calculation of $\partial \mathcal{T}_{st} / \partial \mathbf{c}$ in Eq. (13). In the current implementation, this requires multiple table look-ups due to the presence of the modified basis functions ψ_l^q in Eq. (15), compared to only one for calculating $\partial \mathcal{T}_s / \partial \mathbf{a}$.

III. EXPERIMENTS

Three types of experiments were performed to validate the spatiotemporal deformation model. First, we conducted a prestudy on the temporal dimension of the model separately. Breathing patterns are patient-specific and strong interpatient and intercycle variability has been reported which can affect cycle duration, motion amplitude, and speed of the movement.^{39–41} By fitting the trajectory models to motion data covering many cycles and measured on a large set of patients, we ensured the temporal parametrization is flexible enough to capture the dynamics of respiratory motion.

The most suitable temporal parameter values were retained and used for the spatiotemporal deformation models.

The latter were applied to the registration of 4D CT images of the thorax. Extensive spatial validation of the registrations was performed using large sets of landmarks. The registration accuracy was compared to the conventional frame-to-frame approach. In a final experiment, the benefit of the spatiotemporal approach was illustrated, by assessing the performance in the presence of artifacts.

III.A. Temporal fit of diaphragm motion data

Data description. We used projection sequences of cone-beam computed tomography (CBCT) acquired at the Netherlands Cancer Institute (Amsterdam, the Netherlands) for image-guidance of 33 lung cancer patients treated by radiotherapy with the protocol described by Sonke *et al.*⁴² Cone-beam projections consist of planar x-ray images, acquired from rotating views around the patient. They were acquired at 5.5 fps over 200° with a 50°/min gantry rotation speed for 4D CBCT imaging.⁴³ Projections are 512² matrices with 0.8² mm² spatial resolution (0.52² mm² at the isocenter). 257 sequences of cone-beam projection images (5–19 per patient) were analyzed.

The motion was analyzed by extracting the craniocaudal position of a diaphragm dome using an adapted version⁴¹ of the algorithm developed to extract a respiratory signal for 4D CBCT reconstruction.^{43,44} The extraction resulted in a 2 min 1D+t signal per acquisition with 0.52 mm spatial and 5.5 fps temporal resolutions, i.e., two of the four dimensions of the sought 4D model at fine resolution but for only one point of space. In addition to the largeness of the dataset, the projection images are advantageous because they have higher craniocaudal and temporal resolutions than 4D CT images. As such, the diaphragm motion data provided a valuable benchmark for tuning the temporal parametrization of the deformation models.

Experiments. Each signal was split in respiratory cycles by detecting the end-inhale peaks after smoothing out the local minima. Each cycle was analyzed separately by assuming periodicity, similar to 4D CT images. The temporal models described in Sec. II C were fitted to each cycle with the optimal solution in the least squares sense. The influence of the trajectory model parameters was evaluated: we varied the B-spline order m and the control point spacing s , or equivalently the number of control points. In addition, we verified the suitability of a smoothness constraint at end-inhale by comparing \mathcal{T}_t and \mathcal{T}_t^* .

The similarity between the measured and the fitted signals was evaluated using the root mean square (RMS) of their difference. Results for each patient were averaged and the group mean over all patients was computed. The results were analyzed both globally and per respiratory phase by dividing each cycle into ten equitemporal phase bins, as it is typically done in current 4D CT scanners.

III.B. Spatiotemporal registration of 4D CT

Data description. We used 4D CT data sets of six nonsmall cell lung cancer patients acquired at the Léon Bérard Cancer Center (Lyon, France) for the purpose of radio-

therapy planning on a Philips 16-slice Brilliance Big Bore Oncology Configuration (Phillips Medical Systems, Cleveland, OH). Acquisitions were performed in helical mode using a table pitch of 0.1, 400 mA s effective exposure (80 mA tube current) at 120 kV.

Respiratory-correlated reconstruction was performed through simultaneous acquisition of a respiratory surrogate signal, provided by the Pneumo Chest pressure belt (Lafayette Instrument, Lafayette, IN). Reconstruction yielded ten 3D CT frames at approximately $1 \times 1 \times 2 \text{ mm}^3$ spatial resolution.

Experiments. The spatiotemporal deformable registration approaches \mathcal{T}_{st} and \mathcal{T}_{st}^* , as described in Sec. II D, were applied to all 4D CT data sets. In addition to the normal set of constraints used for \mathcal{T}_t and \mathcal{T}_t^* , constraint (8) was enforced for all registrations. Deformable registration was performed with respect to the middle frame (time point $t_r=5$), roughly corresponding to end-exhale. The position of end-exhale has been reported to be more reproducible than end-inhale,^{42,45} making it a suitable reference to analyze breathing motion.

For validation purposes, anatomical landmarks were identified in the lung region using a semiautomatic software tool.⁴⁶ The system automatically provided a set of well-distributed, distinctive points with index $\mathbf{p}_r \in \mathbf{I}$ in the lung region of the end-exhale frame. Observers identified the corresponding positions $\mathbf{p}_k \in \mathbf{I}$ of the points in frame k , using a custom designed interface and aided by initial estimates provided by the system. Points coinciding with artifacts were excluded. The system initially provided 130 distinctive points and the procedure was stopped after 100 points were successfully identified in the corresponding frames.

For all six patients, 100 point correspondences were provided between the end-exhale and the end-inhale frame, and the process was repeated by a second observer. The mean distance between the annotations was 0.5 mm (0.9 mm standard deviation). For patients 1–3, a single observer provided 100 correspondences for each of the frames of the 4D CT, resulting in a total of 900 manually identified landmarks for each of the three data sets. The manual annotations were compared to the corresponding point positions estimated through registration by computing the target registration error (TRE),

$$\text{TRE} = \|\mathcal{T}_{st}(\mathbf{p}_r, k) - \mathbf{p}_k\|. \quad (16)$$

The registration results were also compared to those obtained when performing consecutive 3D registrations using \mathcal{T}_s , described in Sec. II B. The same multiresolution and optimization scheme was applied as in the spatiotemporal case. Since no temporal regularization is applied in the case of \mathcal{T}_s , the results are considered as a reference indicating the achievable registration accuracy when allowing all temporal degrees of freedom.

III.C. Registration of 4D CT with artifacts

Data description. To dispose of a ground truth, we constructed a 4D CT acquisition with artifacts f_a by introducing a simulated, motion-induced artifact in the 4D CT of patient 2, characterized by large motion. A midinhalation frame ($k=8$) was altered by modifying a series of axial slices halfway the lungs. Ten slices starting from slice index $i_2=i_a$ in the end-exhale frame ($k=5$) were copied to the same location in the target frame, i.e.,

$$f_a(i, k) = \begin{cases} f(i, 5) & \text{for } k=8, \ i_2 \in [i_a, i_a + 10) \\ f(i, k) & \text{otherwise.} \end{cases} \quad (17)$$

The procedure resulted in an axial slab of 20 mm along the craniocaudal direction, containing an inconsistent view of the patient anatomy with respect to the surrounding slices. This resembles the situation of a frame locally influenced by erroneous tagging of the respiratory phase or irregular breathing during image acquisition.

Experiments. The simulated sequence f_a was registered in the same way as described in the previous experiment. The registration results were compared to those obtained using consecutive 3D registrations. By comparing also to the results obtained for the original sequence, the sensitivity of the methods to locally introduced artifacts was evaluated.

The registration accuracy was assessed by using the landmarks identified in the original, unmodified 4D CT acquisition. The analysis is performed at two levels. First, we computed a global evaluation of the TRE, taking into account all

TABLE I. Group mean of the RMS of the difference between the measured motion of the diaphragm dome in the craniocaudal direction and fitted functions for the two temporal models with different number of control points and B-spline degrees. For degrees 0 and 1, both models are equivalent.

Spline degree m	Internal control points						
	4	5	6	7	8	9	10
	Group mean RMS (mm) of \mathcal{T}_t						
0	2.65	2.25	1.95	1.72	1.53	1.39	1.27
1	1.25	0.98	0.83	0.72	0.63	0.57	0.51
2	1.17	0.89	0.77	0.67	0.60	0.54	0.49
3	1.17	0.87	0.76	0.66	0.59	0.53	0.48
	Group mean RMS (mm) of \mathcal{T}_t^*						
2	0.70	0.56	0.47	0.40	0.34	0.30	0.27
3	0.67	0.55	0.47	0.40	0.35	0.30	0.27

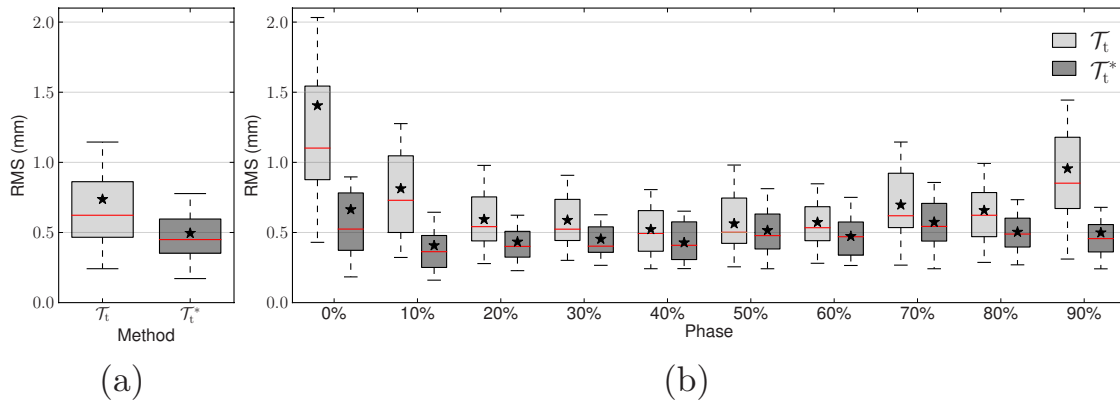


FIG. 2. Box and whisker plots of the RMS errors per patient after fitting the models to the diaphragm motion data, using with five internal control points and $m=3$. (a) The RMS over the entire cycle; (b) the RMS per phase bin.

landmarks. Second, a local analysis was performed by only considering landmarks located within five slices of the artifact.

IV. RESULTS

IV.A. Temporal fit of diaphragm motion data

The fitting results are summarized in Table I for a variety of B-spline functions. Constant B-splines ($m=0$) gave much poorer results than other degrees. This is not surprising, since they produce piecewise constant functions which cannot describe the continuity of the respiratory motion. Linear B-splines ($m=1$) gave residuals of the same order, but were found significantly worse than quadratic ($m=2$) and cubic splines ($m=3$) for all tested models ($p < 3 \times 10^{-4}$). Cubic B-splines consistently gave better results than quadratic splines, although the difference was not significant for all tested models in Table I. As they also inherently impose a stronger temporal regularization, which is our purpose, they were selected for the rest of the study.

The influence of the number of control points can also be seen from Table I. For both models, the residual of the fit was proportional to the spacing of the control points s : the Pearson's product-moment correlation coefficient was

greater than 0.99. As expected, the number of control points is a trade-off between the achievable representation accuracy and the parameters of the fitted function.

We used box and whisker plots to further illustrate the distribution of fitting errors. The box extends from the lower ($p_{25\%}$) to the upper quartile ($p_{75\%}$) of the data, with a horizontal line at the median and a * symbol at the mean. The whiskers extend from the box to the most extreme value below $p_{25\%} + 0.75 \times (p_{75\%} - p_{25\%})$, the remaining points are considered outliers. Outliers were not plotted for clarity.

Figure 2 illustrates the global fitting errors and the errors per respiratory phase bin for \mathcal{T}_t and \mathcal{T}_t^* (using five internal control points and $m=3$). The overall performance of \mathcal{T}_t^* was much better than for \mathcal{T}_t . The largest discrepancies can be observed near end-inhale (0%). At this phase, the change of speed was too sudden to be described by the smooth trajectory model \mathcal{T}_t and the residual was found to be significantly higher ($p < 2 \times 10^{-3}$) than at all other phases for all tested values of m and s . This was not the case for the piecewise smooth \mathcal{T}_t^* model, which resulted in more homogeneous residuals per phase because the smoothness constraint is relaxed at end-inhale.

TABLE II. Summary of the temporal properties for the registration methods when using cubic splines for the spatiotemporal methods, and a control point spacing of either 2 or 2.5 frames. The amount of temporal control points (CPs) reflect the internal CP as well as the ones required at the border. The number of degrees of freedom (DOF) are the number of CP, reduced by the number of constraints. As an example, we list the resulting number of parameters required to register the inner thoracic region (\mathbf{I}_{in}) for patient 1.

Properties	Representation					
	\mathcal{T}_s $s=1$	$s=2$	\mathcal{T}_{st} $s=2.5$	$s=2$	\mathcal{T}_{st}^* $s=2.5$	$s=2.5$
Temporal CP	9	8	7	8	7	
Continuity at $t=0$	/	C^2	C^2	C^0	C^0	
Constraints	0	4	4	2	2	
Temporal DOF	9	4	3	6	5	
Parameters	63 882	28 392	21 294	42 588	35 490	

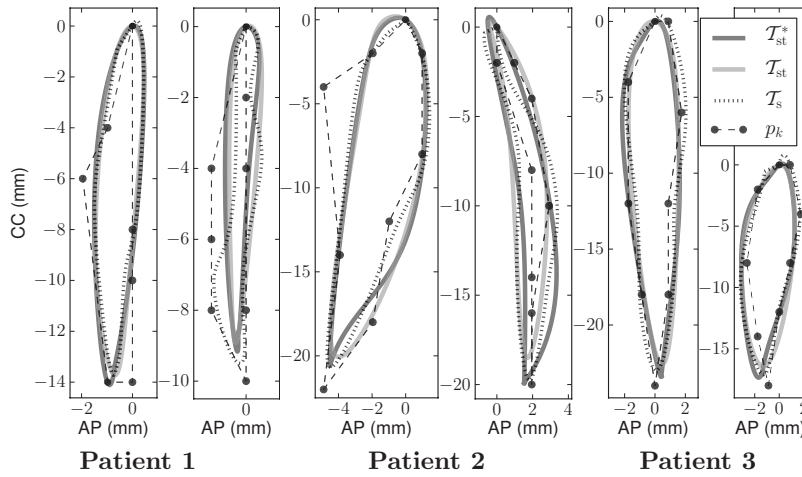


FIG. 3. Trajectories projected on the sagittal plane, for some landmarks with large displacements of patients 1–3. The trajectories obtained using the respective registration methods are plotted along with the manually identified landmark positions throughout the 4D CT (p_k). For all trajectories shown, at least two landmark positions coincided at end-exhale, corresponding to position (0,0).

IV.B. Spatiotemporal registration of 4D CT

We retained the temporal representations with four and five internal control points for the spatiotemporal models, which corresponds to $s=2.5$ and 2 frames, respectively. Table II summarizes the temporal characteristics of the registration methods.

Trajectories obtained for landmarks with large displacements are plotted in Fig. 3. The landmark positions identified manually throughout the 4D CT are also shown and were linearly interpolated for clarity. The estimates obtained using T_s were interpolated using cubic splines. The trajectories of the spatiotemporal methods were directly obtained from the continuous 4D transforms. All trajectories were projected on the sagittal plane, where motion predominantly occurs. Overall, the obtained trajectories appear very similar. The spatiotemporal trajectories tend to be smoother than T_s . The main difference between T_{st} and T_{st}^* is visible at end-inhale (bottom of the plot). At this point, T_{st}^* tends to be pointier and in some cases visibly closer to the corresponding landmark. Note that deviations between the estimated trajectories and the measured landmark trajectories are partially due to the landmark identification process, which was performed in

voxel index space, while the trajectories evolve in the continuous space. This effect will also contribute to the registration errors evaluated using the landmarks.

For patients 1–3, landmarks were available for all frames of the 4D CT. The global registration accuracy is summarized in Table III in terms of the mean TRE based on 900 landmarks each. For spacing $s=2$ frames, the group mean TRE of both spatiotemporal methods was within 0.1 mm of T_s . When increasing the temporal control point spacing from 2 to 2.5, the mean TRE increased gradually (1.27 ± 1.17 mm for T_{st}^* and 1.18 ± 1.03 mm for T_{st}), but remained comparable to T_s . For clarity, only results using a spacing of 2 frames will be shown in the following.

For patients 1–3, the registration errors were also analyzed for each frame separately. Figure 4(a) corresponds to the group mean TRE of the entire 4D CT and Fig. 4(b) shows the group mean TRE per frame. The mean TRE over the entire 4D CT was comparable for all methods, although T_{st} performed slightly worse. The analysis per phase revealed that most discrepancies in TRE are located near end-inhale (0%, 10%, and 90%). T_{st}^* generally obtained an accuracy closer to T_s for these phases.

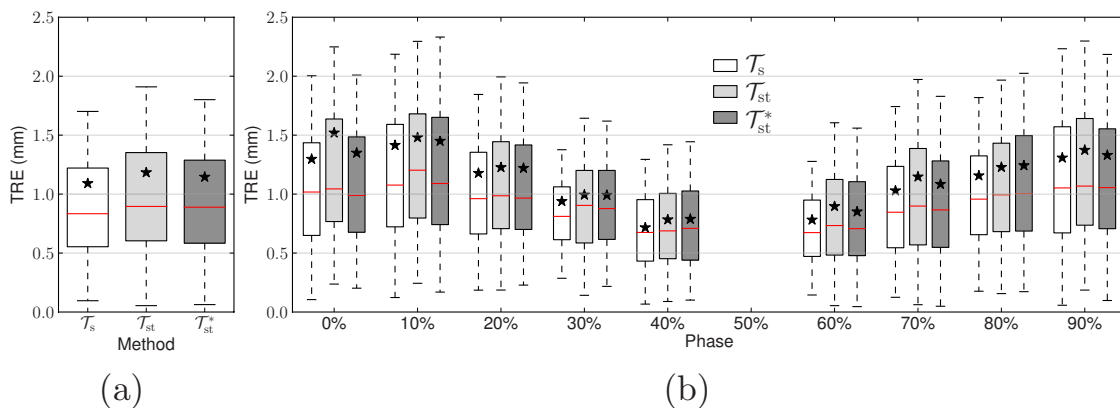


FIG. 4. Box and whisker plots of the group mean TRE for patients 1–3 for which landmarks were available in all frames, using $m=3$ and $s=2$ frames. (a) The combined registration errors for the entire 4D registration. Each box is drawn based on 2700 landmarks. (b) TRE per phase bin. For each frame, the registration error is estimated from 300 landmarks. The 50% phase bin corresponds to the reference frame.

TABLE III. The mean TRE obtained over the nine frames for patients 1–3 based on 900 landmarks each and its group mean (GM). The registration error (± 1 SD) of the 3D registration is compared to the accuracy obtained for the spatiotemporal algorithms with $m=3$ and $s=2$ frames. The original landmark distance (Original) is given to illustrate the magnitude of the motion.

Patient	TRE for 4D CT (mm)			
	Original	\mathcal{T}_s	\mathcal{T}_{st}	\mathcal{T}_{st}^*
1	3.47 ± 2.14	0.96 ± 0.66	1.02 ± 0.71	1.00 ± 0.69
2	6.41 ± 3.99	1.20 ± 0.96	1.37 ± 1.13	1.27 ± 1.09
3	3.65 ± 3.04	1.11 ± 1.14	1.17 ± 1.08	1.16 ± 1.15
GM	4.51 ± 3.15	1.09 ± 0.94	1.19 ± 0.99	1.14 ± 1.00

A separate, more extensive evaluation of the accuracy of the end-exhale to end-inhale registration is listed in Table IV for patients 1–6. \mathcal{T}_{st}^* consistently outperforms \mathcal{T}_{st} in terms of mean TRE. The difference in group mean TRE between \mathcal{T}_{st}^* and \mathcal{T}_s was below 0.1 mm. In contrast, the difference in mean TRE between \mathcal{T}_s and \mathcal{T}_{st} was above 0.1 mm for five out of six patients, and the group mean TRE was almost 0.2 mm higher. This confirms the results reported when fitting the diaphragm motion data in Sec. IV A, where it was found that the smooth temporal model resulted in larger errors near end-inhale.

Table III shows relatively small differences in group mean TRE over the entire 4D CT, suggesting comparable performance for all registration methods. This measure was found misleading, as it tends to average out the differences due to the large numbers of measurements (2700 landmarks for each method). Further analysis showed that the performance of \mathcal{T}_{st} varied considerably from patient to patient. While for patient 1, all methods obtained very similar results, differences in TRE of the order of 0.5 mm were found at certain breathing phases for patient 2 (Fig. 5, note the change in scale with respect to Fig. 4).

TABLE IV. The mean TRE (± 1 SD) obtained by evaluating the registration only at end-inhale for patients 1–6 based on 100 landmarks each, and its GM. The registration error of the 3D registration (\mathcal{T}_i) is compared to the accuracy obtained for the spatiotemporal algorithms (\mathcal{T}_{st} and \mathcal{T}_{st}^*) using $m=3$ and $s=2$ frames. The original landmark distance (Original) is given to illustrate the magnitude of the motion.

Patient	TRE for end-inhale (mm)			
	Original	\mathcal{T}_s	\mathcal{T}_{st}	\mathcal{T}_{st}^*
1	6.34 ± 2.94	0.94 ± 0.51	0.98 ± 0.56	0.96 ± 0.57
2	14.00 ± 7.17	1.44 ± 1.04	1.95 ± 1.88	1.56 ± 1.34
3	7.67 ± 5.03	1.51 ± 1.66	1.63 ± 1.66	1.53 ± 1.70
4	7.33 ± 4.86	1.79 ± 2.71	1.97 ± 3.00	1.96 ± 2.92
5	7.09 ± 5.08	1.43 ± 1.39	1.54 ± 1.49	1.48 ± 1.39
6	6.68 ± 3.67	1.18 ± 0.80	1.32 ± 1.13	1.25 ± 0.95
GM	8.19 ± 4.97	1.38 ± 1.53	1.57 ± 1.78	1.46 ± 1.65

TABLE V. The mean TRE for the 4D CT sequence with simulated artifacts (f_a) and for the original, unmodified 4D CT (f) corresponding to patient 2. The evaluation is limited to the frame where the artifact is introduced. The global TRE is based on 100 landmarks. The local TRE is based on 24 landmarks, all within five slices of the inserted artifact. The registration error of the 3D registration (\mathcal{T}_s) is compared to the accuracy obtained for the spatiotemporal algorithm (\mathcal{T}_{st}^* using $m=3$ and $s=2$ frames). The original landmark distance (Original) is given to illustrate the magnitude of the motion.

Measure	Data	TRE (mm)		
		Original	\mathcal{T}_s	\mathcal{T}_{st}^*
Global	f	9.00 ± 3.93	1.42 ± 1.30	1.44 ± 1.16
	f_a		3.17 ± 3.47	1.57 ± 1.20
Local	f	11.40 ± 3.74	1.38 ± 1.44	1.46 ± 1.05
	f_a		6.82 ± 4.38	1.90 ± 1.22

IV.C. Spatiotemporal registration of 4D CT with artifacts

The registration accuracy obtained for the sequence f_a is summarized in Table V. We only report results using the piecewise smooth spatiotemporal model \mathcal{T}_{st}^* . We also list the TRE obtained for the original 4D CT, corresponding to patient 2. With respect to the original 4D CT, the local and global TREs of \mathcal{T}_s and \mathcal{T}_{st}^* are within 0.1 mm. After inserting the artifact, the global TRE more than doubles for \mathcal{T}_s , while the TRE of the spatiotemporal method increases only marginally. Locally, the influence of the artifact is even more noticeable for \mathcal{T}_s . For the spatiotemporal approach, however, the local TRE remains below 2 mm.

Figure 6 shows the motion fields obtained using \mathcal{T}_s and \mathcal{T}_{st}^* for the inner thoracic region. The top row corresponds to the original 4D CT acquisition of patient 2. Both methods produce very similar motion fields. The main differences can be observed near the diaphragm. The bottom row corresponds to the sequence with artifacts f_a . In this case, strong perturbations can be observed in the motion field obtained using \mathcal{T}_s . The influence of the introduced artifact is also noticeable for \mathcal{T}_{st}^* , but the changes in the motion field are less dramatic.

In Fig. 7, three examples are given of motion fields in the presence of real artifacts. The artifacts are shown in the first column and are generally most clearly visible near the diaphragm. The second column shows the motion fields obtained using \mathcal{T}_s . Strong perturbations can be seen, mainly in the part of the motion field that maps to the region of the artifact; i.e., slightly above the location of the artifacts. The spatiotemporal approach tends to be less influenced by the artifacts. The resulting motion fields are noticeably smoother making them more plausible from a physiological point of view.

V. DISCUSSION

Temporal constraints. In Sec. IV A, the smooth and piecewise smooth temporal models were compared at equal control point spacing (see Fig. 2). The difference in temporal constraints between \mathcal{T}_i and \mathcal{T}_i^* results in a different number of

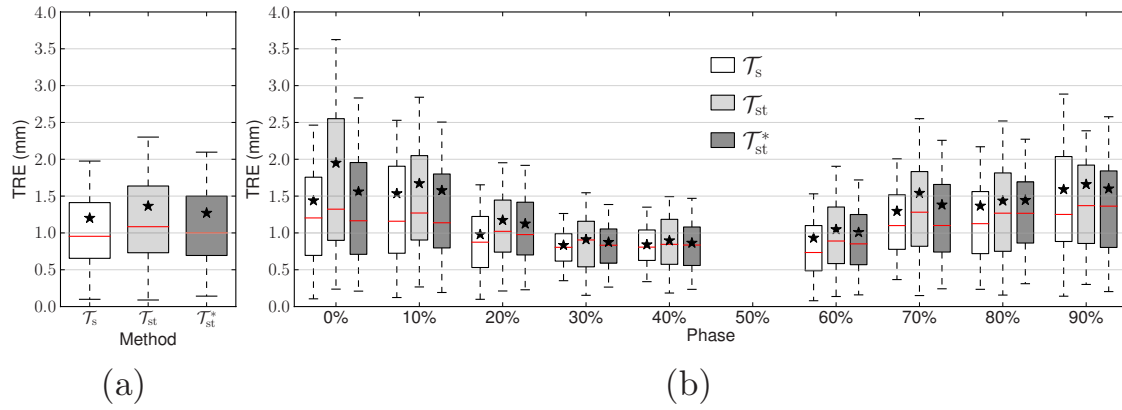


FIG. 5. Box and whisker plots of TRE for patients 2 for which landmarks were available in all frames, using $m=3$ and $s=2$ frames. (a) The combined registration errors for the entire 4D registration. Each box is drawn based on 900 landmarks. (b) TRE per phase bin. For each frame, the registration error is estimated from 100 landmarks. The 50% phase bin corresponds to the reference frame.

degrees of freedom at equal control point spacing. We therefore also performed a comparison between both models at equal degrees of freedom. The global performance of \mathcal{T}_l was still significantly worse ($p < 4 \times 10^{-2}$) compared to the corresponding \mathcal{T}_l^* models (see Table I for \mathcal{T}_l^* with two control points less than \mathcal{T}_l). In addition, despite the *global* increase in degrees of freedom, the high fitting residual at end-inhale remained for \mathcal{T}_l .

This confirms that a *local* increase in control points (as is the case for the piecewise smooth model \mathcal{T}_l^*) is more efficient in terms of number of parameters, to obtain an accurate representation throughout the respiratory cycle. It should not be excluded that other piecewise models can be found, requiring less degrees of freedom, while obtaining a similar accuracy. These could consist in making sensible assumptions about the trajectory near end-inhale that can be translated into con-

straints, eliminating one or both of the additional degrees of freedom.

The end-exhale phase, although also characterized with inversion of the motion, did not require further investigation of the constraints. It has been reported that respiratory motion tends to be asymmetrical,^{45,47} spending more time near end-exhale than end-inhale. Phase bins near end-exhale will represent relatively small deformation with respect to each other. Uniformly spaced control points with respect to these bins will lead to a spatially higher control point density near end-exhale, allowing a more accurate representation, even in the presence of smoothness constraints. This is confirmed by

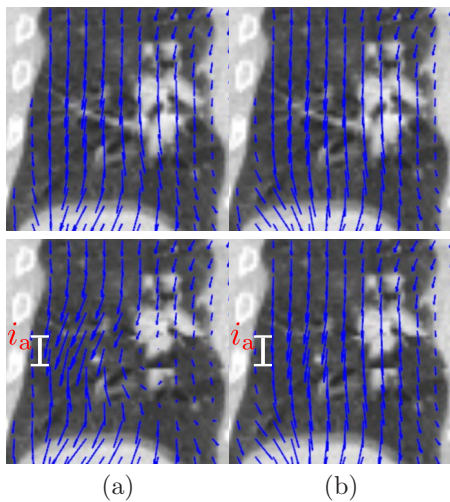


FIG. 6. Motion fields in the presence of simulated artifacts: the top row corresponds to the original 4D CT acquisition of patient 2, the bottom row corresponds to the modified sequence f_a in which an artifact was inserted at position i_a . (a) Coronal view of the motion field obtained for \mathcal{T}_s and (b) corresponding view of the motion field obtained for \mathcal{T}_{st}^* using $m=3$ and $s=2$ frames.

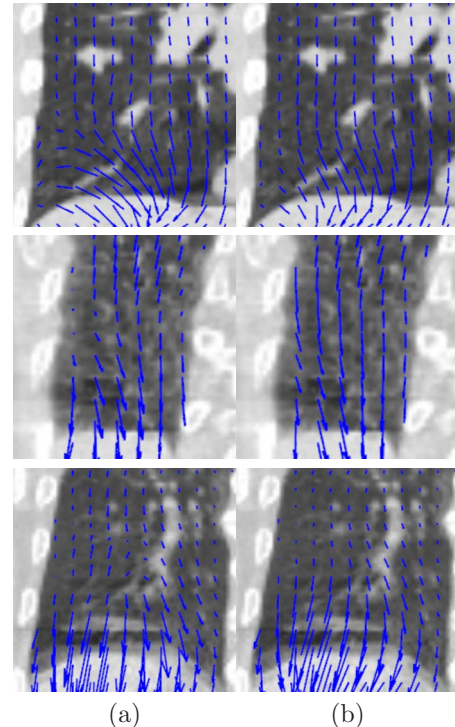


FIG. 7. Three examples of motion fields in the presence of real artifacts. (a) Motion fields obtained for \mathcal{T}_s and (b) \mathcal{T}_{st}^* using $m=3$ and $s=2$ frames.

the trajectories depicted in Fig. 3, where two to three landmarks coincided with the end-exhale position.

From a temporal constraints point of view, \mathcal{T}_t^* is related to the trajectory model developed by Castillo *et al.*,²⁹ where a compressible flow algorithm is extended with local trajectory modeling to perform 4D motion estimation for 4D CT. In this case, however, one-way (and not cyclic) trajectories were sought between end-inhale and end-exhale, thus not requiring further attention at end-inhale. Cubic polynomials (equivalent to four degrees of freedom) were found to provide sufficient flexibility to parametrize the sought trajectories spanning six frames of the 4D CT. This corresponds well to the six degrees of freedom (\mathcal{T}_{st}^* with a temporal control point spacing of 2) describing the trajectory over 10 frames.

Spatiotemporal registration. The principal aim of this study was to develop a low-dimensional spatiotemporal deformation model to improve robustness of the subsequent registration. We pursued a restrictive parametrization and strong temporal regularization, as these were expected to reduce sensitivity to noise and artifacts. The parametrization was thoroughly investigated, both spatially and temporally, to ensure that an accurate representation of breathing motion was maintained.

Based on the fitting experiments of the diaphragm motion data, cubic temporal B-splines were found to perform best and were selected for the temporal parametrization. The value of the temporal control point spacing s was found to represent a trade-off between achievable accuracy on one hand and an increase of parameters on the other, the latter likely to increase sensitivity to noise and artifacts. In practice, its value should reflect the needs of the application and the quality of the images. Using $s=2$ frames, the spatiotemporal models obtained results comparable to the reference \mathcal{T}_s method and was considered a suitable compromise for the 4D CT images dealt with in this study.

From a parametrization point of view, the \mathcal{T}_{st} model represents interesting characteristics. Minimal curvature is enforced throughout the entire cycle, and about a third less parameters are required with respect to \mathcal{T}_{st}^* . Unfortunately, detailed analysis revealed larger TRE near end-inhale for \mathcal{T}_{st} , indicating the smooth model fails to capture the full extent of the motion. Even though trajectories are expected to be smooth functions of time, a temporally smooth parametrization was found to provide a less accurate representation due to the low temporal resolution of respiratory-correlated imaging and the control point grid. Using \mathcal{T}_{st}^* a uniform performance over the breathing cycle was obtained for all patients, and the group mean TRE was within 0.1 mm of the reference \mathcal{T}_s , for both $s=2$ and 2.5 frames.

The improved matching of \mathcal{T}_{st}^* at end-inhale comes at the price of two additional temporal degrees of freedom with respect to the smooth model. Nonetheless, with respect to \mathcal{T}_s , this model reduces the number of parameters to be estimated during registration by 33% and 45% for $s=2$ and 2.5 frames, respectively (Table II). The impact of this compact, spatiotemporal parametrization was illustrated in the experiment in which a simulated motion-induced artifact was introduced in a 4D CT sequence. The motion field obtained using spa-

tiotemporal registration was found to be considerably less influenced by the artifact, in comparison to the result obtained using \mathcal{T}_s .

Influence of binning and tagging. In Sec. II, we made the assumption that the fourth image dimension was time. This allowed us to interpret \mathcal{T}_t as a trajectory in function of time and its derivatives as velocity and acceleration. In the case of 4D CT imaging, each frame is composed of data acquired at different times and different table positions. The interpretation of the fourth image dimension is closely related to the binning of acquired data, which is usually based on a surrogate signal.

The 4D CT data presented here were obtained from phase-based binning, which is by far the most common procedure. For each acquired cycle, the reconstruction of the different frames is performed by selecting projections equally spaced in time. Ignoring the nonperiodic nature of breathing motion, the images obtained in this fashion can be considered equivalent to a temporal sequence. Alternative binning criteria^{48,49} have been proposed. In particular, amplitude-based binning is expected to provide frames uniformly spaced with respect to organ displacement. For such images, other smoothness constraints at end-inhale and even end-exhale might be more suitable.

For all data presented, individual cycles were detected by providing tags at end-inhale. The diaphragm motion data were artificially made periodic at end-inhale, to allow fitting the cyclic trajectory models. This procedure can be held partly responsible for the rapid changes at end-inhale and contributes to the fitting residuals. To quantify the effect, we repeated the experiments when tagging at end-exhale. Comparatively larger residuals were observed at end-exhale, indicating the influence of the tagging position. The highest residuals were, however, still observed at end-inhale, confirming they are indeed caused by the sudden change in motion.

Robustness to artifacts. Thoroughly evaluating the performance in the presence of artifacts is difficult due to the absence of a ground truth for the underlying image. The improved robustness of the spatiotemporal approach was therefore illustrated using a simple experiment based on simulated data and through visual inspection of motion fields for real artifacts. Further analysis of the behavior of the spatiotemporal model in the presence of artifacts or noise is required. In particular, the influence of the temporal size and location of artifacts merits attention.

In the 3D, Eq. (2), and 4D criteria, Eq. (12), regularization terms penalizing undesirable properties of the deformation field were not included. The present study was limited to investigating the impact of explicit parametric restrictions. Regularization penalties will provide additional robustness and are expected to be complementary to the parametric contributions. In addition, the spatiotemporal framework allows regularization schemes to be extended to the temporal dimension, as in Ref. 50.

Applications of spatiotemporal motion analysis. Spatiotemporal deformation model (11) was applied to 4D CT of the thorax in a 3D-4D registration framework. The model

can also be applied to spatiotemporal motion analysis between sequences. By relaxing condition (8) which constrains the deformation at the reference frame, the model can be applied to a 4D-4D, frame-to-frame registration framework. By replacing criterion (12) with a suitable similarity measure, other modalities and even multimodal problems can be studied. This is similar to the 4D-4D registration method for respiratory-correlated images described by Schreibmann *et al.*²⁸ or the spatiotemporal alignment of cardiac sequences presented by Perperidis *et al.*²¹ In comparison, our method assumes only frame-to-frame spatial deformations without temporal shifts and constrains deformations to a cyclic trajectory. These assumptions limit the degrees of freedom and should be well suited to analyze motion patterns between, for instance, respiratory-correlated 4D CT, 4D CBCT (Ref. 43) or 4D magnetic resonance images.⁵¹

VI. CONCLUSION

We developed a spatiotemporal deformation model for deformable registration of respiratory-correlated images of the thorax. The model was obtained by extending spatial free-form deformations to the temporal domain, using a cyclic trajectory model based on cubic B-splines. A piecewise smooth temporal parametrization was found most suitable to account for the rapid changes in velocity at end-inhale. The spatiotemporal modeling resulted in a considerably more compact description of the deformation model. Spatiotemporal registration leads to comparable registration results while improving the robustness to artifacts.

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