# Splines for diffeomorphisms 

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#### Abstract

This paper develops a method for higher order parametric regression on diffeomorphisms for image regression. We present a principled way to define curves with nonzero acceleration and nonzero jerk. This work extends methods based on geodesics which have been developed during the last decade for computational anatomy in the large deformation diffeomorphic image analysis framework. In contrast to previously proposed methods to capture image changes over time, such as geodesic regression, the proposed method can capture more complex spatio-temporal deformations.

We take a variational approach that is governed by an underlying energy formulation, which respects the nonflat geometry of diffeomorphisms. Such an approach of minimal energy curve estimation also provides a physical analogy to particle motion under a varying force field. This gives rise to the notion of the quadratic, the cubic and the piecewise cubic splines on the manifold of diffeomorphisms. The variational formulation of splines also allows for the use of temporal control points to control spline behavior. This necessitates the development of a shooting formulation for splines.

The initial conditions of our proposed shooting polynomial paths in diffeomorphisms are analogous to the Euclidean polynomial coefficients. We experimentally demonstrate the effectiveness of using the parametric curves both for synthesizing polynomial paths and for regression of imaging data. The performance of the method is compared to geodesic regression.


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## 1. Introduction

With the now common availability of longitudinal and time series image data, models for their analysis are critically needed. In particular, spatial correspondences need to be established through image registration for many medical image analysis tasks. While this can be accomplished by pair-wise image registration to a template image, such an approach neglects spatio-temporal data aspects. Instead, explicitly accounting for spatial and temporal dependencies is desirable.

A common way to describe differences in geometry of objects in images is to summarize them using transformations. Transformations are fundamental mathematical objects and have long been known to effectively represent biological changes in organisms (Amit et al., 1991; Thompson et al., 1942). The field of computational anatomy (Grenander and Miller, 1998; Miller, 2004; Miller et al., 1997; Thompson and Toga, 2002) provides a rich mathematical setting for statistical analysis of complex geometrical structures seen in 3D medical images. At its core, computational anatomy is based on the representation of anatomical shape and its variability using smooth and

[^0]invertible transformations that are elements of the nonflat manifold of diffeomorphisms with an associated Riemannian structure. The large deformation diffeomorphic metric mapping (LDDMM) framework of computational anatomy exploits ideas from fluid mechanics and builds maps of diffeomorphisms as flows of smooth velocity fields (Younes, 2010; Younes et al., 2009).

Research in the last decade provideds several methods to represent natural biological variability by modeling them as nonlinear transformations in the manifold of diffeomorphisms. Their focus has primarily been on geodesic models. For example, methods of Fréchet mean (Davis, 2008), geodesic regression (Niethammer et al., 2011) and hierarchical geodesic models (Singh et al., 2013a) are first order models that rely on computing geodesics within the space of diffeomorphisms. While such models have proven to be effective, their use is limited to modeling only "geodesic-like" image data. However, geodesics are not always appropriate for regression modeling of time series data. In particular, nonmonotonous shape changes seen in time sequence or videos of medical images of periodic breathing, cardiac motion, or shape changes in the human brain during a long age range (10-90 years), do generally not adhere to constraints of geodesicity. This necessitates the development of higher order models of regression within the space of diffeomorphic transformations. Computational anatomy has seen very little work on


Fig. 1. Models of parametric regression for computational anatomy. Geodesic regression generalizes the notion of parametric linear regression in the Euclidean spaces (left) to the group of diffeomorphisms (right). The model estimate comprises of the initial velocity, $v(0)$, at the identify diffeomorphism, $e$, and completely parameterizes the best fit regression geodesic path, $\phi(t)$. However, no such generalizations of the known Euclidean models of higher order parametric regression such as the cubics or the splines (left) exist for the group of diffeomorphisms (right).
higher-order models of registrations for modeling image time series (Fig. 1).

Contribution. In this article we propose:

1. an acceleration-controlled model that generalizes the idea of cubic curves to manifold of diffeomorphisms and is capable of modeling nonmonotonic shape changes under the large deformation (LDDMM) setting,
2. a shooting based solution to cubic curves that enables parametrization of the full regression path using only initial conditions,
3. a method of shooting cubic splines as smooth curves to fit complex shape trends while keeping data-independent (finite and few) parameters, and
4. a numerically practical algorithm for regression of "non-geodesic" medical imaging data.

The work described in this manuscript significantly extends our work presented at MICCAI (Singh and Niethammer (2014)). In particular, (1) we make use of a new formulation directly advecting the inverse of a diffeomorphism, (2) we provide extended discussions of the approach, and (3) present a variety of new results to illustrate the behavior of the approach.

### 1.1. Related work

Methods that generalize Euclidean parametric regression models to manifolds have proven to be effective for modeling the dynamics of changes represented in time series of medical images. For instance, methods of geodesic image regression (Niethammer et al., 2011; Singh et al., 2013b) and longitudinal models on images (Singh et al., 2013a) generalize linear and hierarchical linear models, respectively. Although the idea of polynomials (Hinkle et al., 2014) and splines (Trouvé \& Vialard, 2012) on the landmark representation of shapes has been proposed, higher-order extensions for image regression remain deficient. While Hinkle et al. (2014) develop an approach for general polynomial regression and demonstrate it on finite-dimensional Lie groups, infinite dimensional regression is demonstrated only for first-order geodesic image regression.

These parametric regression models are advantageous since their estimated parameters can be used for further statistical analysis. For instance, initial momenta obtained from Fréchet atlas construction of a population of images can be treated as signature representations of shape differences across the group and can be treated as features to train classification and regression models (Singh et al., 2014).

Machado et al. (2006) also discuss the notion of first order variational fitting of curves to data on Riemannian manifolds. The solution to the variational problem results in piecewise geodesics, where the number of pieces is equal to the number of manifold data points. Durrleman et al. (2013) present a method of regression analysis of population of time series of shapes based on timewarping. This model
is presented for regression of shapes that do not trivially generalize to the regression of image time series data. The model also results in piecewise geodesics to summarize individual spatiotemporal trends such that the number of pieces is data dependent. Following the earliest ideas presented in Noakes et al. (1989), Camarinha et al. (1995), Crouch and Leite (1995), the work of Machado et al. (2010) further develops the notion of variational fitting to estimating piecewise higher order curves on data in general Riemannian manifolds. The solution involves estimating the Riemannian curvature tensor and is useful for finite dimensional manifolds where the tensor can be evaluated analytically. Krakowski (2003) provides a theoretical review of variational splines and explores, in particular, splines on finite-dimensional manifolds such as the space of rotations, $\mathbb{S O}(3)$, and the unit sphere, $\mathbb{S}^{n}$.

Other regression methods include those by Davis et al. (2010), Lorenzi et al. (2010), Vercauteren et al. (2009), Schwartz et al. (2015) and Gu et al. (2006). Davis et al. (2010) generalize the notion of kernel regression to manifolds. Kernel regression is a nonparametric approach and hence does not provide a summary representation of the regression fit in terms of a finite set of parameters for further analysis. Lorenzi et al. (2010) propose a smooth spatiotemporal modeling of image time series data using the regression of pairwise registrations under the stationary velocity field (SVF) framework of LogDemons (Vercauteren et al., 2009). More recently, Schwartz et al. (2015) extend this idea and propose locally linear regression under the SVF framework. Gu et al. (2006) develop the spline interpolation for the case when the domain of the independent variable itself is a manifold. This is useful for surface interpolation, for example, in graphics or geometric design to interpolate surfaces represented on manifold domains that give rise to shapes with arbitrary topologies.

Relevant background readings include those by Noakes et al. (1989), Camarinha et al. (1995), Crouch and Leite (1995), where the notion of splines on general Riemannian manifolds were first introduced. These series of papers discuss a theoretical characterization of the general variational cubic curves and spline interpolation. In this paper, we present a shooting based formulation of the classical variational spline formulation and derive its solution and numerical estimation procedures for the group of diffeomorphisms.

The remainder of this article is structured as follows: Section 2 reviews the variational approach to splines in Euclidean space and motivate its shooting formulation for parametric regression. Section 3 then generalizes this concept of shooting splines for diffeomorphic image regression. We discuss experimental results in Section 4, and conclude the article with a discussion of future work in Section 5 .

## 2. Shooting-splines in the Euclidean case

To motivate our formulation for splines on diffeomorphisms it is instructive to first revisit the variational formulation for splines in the Euclidean case. This facilitates a more straightforward presentation

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of the fundamental approach and allows to make direct connections to the formulation for splines on diffeomorphisms.

### 2.1. Variational formulation

An acceleration controlled curve with time-dependent states, $\left(x_{1}\right.$, $\left.x_{2}, x_{3}\right)$ such that, $\dot{x}_{1}=x_{2}$ and $\dot{x}_{2}=x_{3}$, defines a cubic curve in Euclidean spaces for a constant acceleration, $x_{3}$. Here, $x_{1}$ denotes position and $x_{2}$ velocity. In particular, such a cubic curve minimizes an energy of the form, $E=\frac{1}{2} \int_{0}^{1}\left\|x_{3}\right\|^{2} d t$ subject to the dynamic constraints above. The corresponding constrained optimization problem can be written as

$$
\begin{equation*}
\operatorname{minimize}_{x_{1}, x_{2}, x_{3}} E\left(x_{3}\right) \text { subject to } \quad x_{2}=\dot{x}_{1} \quad \text { and } \quad x_{3}=\dot{x}_{2} \tag{1}
\end{equation*}
$$

Here $x_{3}$ is referred to as the control variable that describes the acceleration of the dynamics in this system. The unconstrained Lagrangian for the above is,

$$
\begin{aligned}
E\left(x_{1}, x_{2}, x_{3}, \mu_{1}, \mu_{2}\right)= & \frac{1}{2} \int_{0}^{1}\left\|x_{3}\right\|^{2} d t+\int_{0}^{1} \mu_{1}^{T}\left(\dot{x}_{1}-x_{2}\right) d t \\
& +\int_{0}^{1} \mu_{2}^{T}\left(\dot{x}_{2}-x_{3}\right) d t
\end{aligned}
$$

where $\mu_{1}$ and $\mu_{2}$ are the time-dependent Lagrangian variables or the adjoint variables (also called duals) that enforce the dynamic constraints. Optimality conditions on the gradients of the above Lagrangian with respect to the states, $\left(x_{1}, x_{2}, x_{3}\right)$, result in the adjoint system of equations, $\dot{\mu}_{1}=0$ and $\dot{x}_{3}=-\mu_{1}\left(\mu_{2}\right.$ gets eliminated). This allows for a relaxation solution to Eq. (1), where the state of the system is the full time-course of states, i.e., $\left(x_{1}(t), x_{2}(t), x_{3}(t)\right)$, and the condition $x_{3}(t)=$ const. will be fulfilled at convergence. However, we may also formulate this problem with respect to initial conditions alone, amounting to a shooting solution as discussed in Section 2.2.

### 2.2. From relaxation to shooting

A relaxation solution has originally been proposed for diffeomorphic image registration by Beg et al. (2005). Here, a full-spatiotemporal velocity field was the variable to be estimated. Instead, a shooting reformulation (Vialard et al., 2011) allowed to represent the image registration problems by optimizing over an initial image and an initial momentum. In the scalar-valued setting the shootingformulation corresponds to optimizing over the initial $y$-intercept and slope of a line, thereby searching over the space of straight lines instead of converging to a straight line as in the relaxation setting. Shooting thereby allowed the formulation of geodesic regression approaches (Niethammer et al., 2011) where one aims to determine the best geodesic fitting the given data and optimized over the initial conditions specifying the geodesic only. Hence, to allow for splines on diffeomorphisms we also need a shooting formulation to be able to compactly represent splines and to express the equivalent of piecewise cubic curves. In the scalar-valued case such a shooting formulation can be obtained by explicitly adding the evolution of $x_{3}$, obtained by solving the relaxation problem, as a dynamical constraint. This increases the order of the dynamics. Denoting, $x_{4}=-\mu_{1}$, results in the classical system of equations for shooting cubic curves,
$\dot{x}_{1}=x_{2}(t), \quad \dot{x}_{2}=x_{3}(t), \quad \dot{x}_{3}=x_{4}(t), \quad \dot{x}_{4}=0$.
The states, $\left(x_{1}, x_{2}, x_{3}, x_{4}\right)$, at all times, are entirely determined by their initial values $\left(x_{1}^{0}, x_{2}^{0}, x_{3}^{0}, x_{4}^{0}\right)$, and in particular, $x_{1}(t)=x_{1}^{0}+x_{2}^{0} t+$ $\frac{x_{3}^{0}}{2} t^{2}+\frac{x_{4}^{0}}{6} t^{3}$. Also note that $x_{4}$ is the derivative of acceleration, $x_{3}$, and can therefore be interpreted as jerk. For a cubic the jerk is constant. When piecing together multiple cubic curves, as will be described in Section 2.3, the jerk will be allowed to jump.


Fig. 2. Gluing together cubics to construct a piecewise cubic curve. This example uses three partitions, $\mathcal{I}_{1}, \mathcal{I}_{2}$ and $\mathcal{I}_{3}$ defined by placing controls at two locations $(C=2)$.

### 2.3. Shooting-splines with data-independent controls for regression

We now present our proposed method of regression using cubic splines using the shooting equations. The goal is to define a smooth curve that best fits the data in the least-squares sense. Since a cubic polynomial by itself is restricted to only fit "cubic-like" data, we propose to add flexibility to the curve by piecing together piecewise cubic polynomials. In other words, we define controls at pre-decided locations in time where the state $x_{4}$ is allowed to jump.

Let, $y_{i}$, for $i=1 \ldots N$, denote $N$ measurements at timepoints, $t_{i} \in$ $(0,1)$. Let $t_{c} \in(0,1)$, for $c=1 \ldots C$, denote $C$ data-independent fixed control locations. For notational convenience, we assume there are no measurements at the end points, $\{0,1\}$, or at the control locations, $\left\{t_{c}\right\}$. The control locations also implicitly define $C+1$ intervals or partitions in $(0,1)$. Let us denote these intervals as $\mathcal{I}_{c}$, for $c=1 \ldots(C+1)$. In particular, since we assume no measurements at the end points and at control locations, it is convenient to define each interval as an open interval, such that, $\mathcal{I}_{1}:=\left(0, t_{1}\right), \mathcal{I}_{2}:=\left(t_{1}, t_{2}\right), \ldots, \mathcal{I}_{c}:=\left(t_{c-1}, t_{c}\right), \ldots$, $\mathcal{I}_{C+1}:=\left(t_{C}, 1\right)$. The constrained energy minimization that solves the regression problem with such a data configuration can be written as,

$$
\operatorname{minimize}_{x_{2}(0), x_{3}(0), x_{4}(0),\left\{x_{4}\left(t_{c}\right)\right\}} \frac{1}{2 \sigma^{2}} \sum_{c=1}^{C+1} \sum_{i \in \mathcal{I}_{c}}\left\|x_{1}\left(t_{i}\right)-y_{i}\right\|^{2}
$$

$$
\text { s.t. } \quad \dot{x}_{1}=x_{2}(t), \quad \dot{x}_{2}=x_{3}(t), \quad \dot{x}_{3}=x_{4}(t), \quad \dot{x}_{4}=0,
$$

(within each interval, $\mathcal{I}_{c}$ ), and
s.t. $\quad x_{1}, x_{2}$, and $x_{3}$ are continuous across $C$.

The partitioning of the domain of independent variable for regression for the case of three partitions using two control locations is depicted in Fig. 2.

The unconstrained Lagrangian enforcing shooting and continuity constraints using time-dependent adjoint states, ( $\lambda_{1}, \lambda_{2}, \lambda_{3}, \lambda_{4}$ ), and duals, ( $\nu_{1}, \nu_{2}, \nu_{3}$ ), is

$$
\begin{aligned}
& E\left(x_{1}^{0}, x_{2}^{0}, x_{3}^{0}, x_{4}^{0}, x_{4}^{t_{c}}, \lambda_{1}, \lambda_{2}, \lambda_{3}, \lambda_{4}, v_{1}, \nu_{2}, \nu_{3}\right)= \\
& \frac{1}{2 \sigma^{2}} \sum_{c=1}^{C+1} \sum_{i \in \mathcal{I}_{c}}\left\|x_{1}\left(t_{i}\right)-y_{i}\right\|^{2} \\
& \quad+\int_{0}^{1}\left(\lambda_{1}^{T}\left(\dot{x}_{1}-x_{2}\right)+\lambda_{2}^{T}\left(\dot{x}_{2}-x_{3}\right)+\lambda_{3}^{T}\left(\dot{x}_{3}-x_{4}\right)\right. \\
& \left.\quad+\lambda_{4}^{T} \dot{x}_{4}\right) d t+v_{1}\left(x_{1}^{-}\left(t_{c}\right)-x_{1}^{+}\left(t_{c}\right)\right) \\
& \quad+v_{2}\left(x_{2}^{-}\left(t_{c}\right)-x_{2}^{+}\left(t_{c}\right)\right)+v_{3}\left(x_{3}^{-}\left(t_{c}\right)-x_{3}^{+}\left(t_{c}\right)\right) .
\end{aligned}
$$

The gradients of this Lagrangian with respect to the adjoint variables result in the forward dynamical constraints. However, the gradients with respect to the primal variables at the optimum result in the adjoint system of equations, $\dot{\lambda}_{1}=0, \dot{\lambda}_{2}=-\lambda_{1}, \dot{\lambda}_{3}=-\lambda_{2}, \dot{\lambda}_{4}=-\lambda_{3}$. The gradients with respect to the initial conditions for states $x_{1}^{0}$ for $l=1, \ldots, 4$ are, $\delta_{x_{1}^{0}} E=-\lambda_{1}(0), \delta_{x_{2}^{0}} E=-\lambda_{2}(0), \delta_{x_{3}^{0}} E=-\lambda_{3}(0)$ and $\delta_{x_{4}^{0}} E=-\lambda_{4}(0)$. The jerks at controls, $x_{4}^{t_{c}}$, are updated using, $\delta_{x_{c}} E=-\lambda_{4}\left(t_{c}^{+}\right)$. The values of adjoint variables required in these gradients are computed by integrating backward the adjoint system.

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Fig. 3. States for splines regression in Euclidean space with one control at $t=0.5$.

Note that $\lambda_{1}, \lambda_{2}$ and $\lambda_{3}$ are continuous at joins, but $\lambda_{1}$ jumps at the data-point location as per, $\lambda_{1}\left(t_{i}^{+}\right)-\lambda_{1}\left(t_{i}^{-}\right)=\frac{1}{\sigma^{2}}\left(x_{1}\left(t_{i}\right)-y_{i}\right)$. During backward integration, $\lambda_{4}$ starts from zero at each interval at $t_{c+1}$ and the accumulated value at $t_{c}$ is used for the gradient update of $x_{4}\left(t_{c}\right)$.

It is critical to note that, along the time, $t$, such a formulation guarantees that: (a) $x_{4}(t)$ is piecewise constant, (b) $x_{3}(t)$ is piecewise linear, (c) $x_{2}(t)$ is piecewise quadratic, and (d) $x_{1}(t)$ is piecewise cubic. Thus, this results in a cubic-spline curve. Fig. 3 demonstrates this shooting spline fitting on scalar data. While it is not possible to explain this data with a simple cubic curve alone, it suffices to allow one control location to recover the meaningful underlying trend. The state, $x_{4}$, experiences a jump at the control location that integrates up thrice to give a $C^{2}$-continuous evolution for the state, $x_{1}$.

## 3. Shooting-splines for diffeomorphisms

Our goal is to generalize the variational approach to cubic splines to the group of diffeomorphism to define splines that can capture for example complex image deformations over time. Our approach is completely analogous to the scalar-valued case described in earlier sections. Before presenting the formulation of cubic curves and splines for diffeomorphisms we briefly review the necessary background in the next section.

### 3.1. Background on the Lie group of diffeomorphisms

Diffeomorphisms offer a way to represent smooth and invertible spatial transformations that match one shape to another. For the purpose of this paper, the shapes refer to objects embedded in 2D or 3D images. We define an image, $I$, as a real-valued $L^{2}$ function on a domain $\Omega \subset \mathbb{R}^{d}$, where $d=2$ or $d=3$ for 2D or 3D images, respectively.

We define a diffeomorphism $g$ as a mapping of $\Omega$ that assigns every point $x \in \Omega$ a new position $x^{\prime}=\phi(x) \in \Omega$. Under this definition, we restrict to transformations that satisfy the following rules of smooth bijection, $g$ should be:

1. Onto: All points in $x^{\prime} \in \Omega$ should be image of some point in domain $\Omega$.
2. One-to-one: Two different points should not map to one single point, i.e., $g(x)=g(y) \Longleftrightarrow x=y$.
3. Smooth: $g$ is $C^{\infty}$ or more generally $C^{k}$, i.e., $k$ differentiable.
4. Smooth inverse: $\phi^{-1}$ is $C^{\infty}$ or more generally $C^{k}$, i.e., $k$ differentiable.

The deformation of an image $I$ by $g$ is defined as the action of the diffeomorphism, given by $g \cdot I=I \circ g^{-1}$. A natural way for generating diffeomorphic transformations is by the integration of ordinary differential equations (ODE) on $\Omega$ defined via the smooth time-indexed velocity vector fields $v(t, x):(t \in[0,1], x \in \Omega) \rightarrow \mathbb{R}^{3}$. The function
$g^{v}(t, x)$ given by the solution of the ODE $\frac{d y}{d t}=v(t, y)$ with the initial condition $y(0)=x$ defines a diffeomorphism of $\Omega$. Diffeomorphisms thus generated as flows of velocity fields form a group under the composition operation and are denoted by $\operatorname{Diff}(\Omega)$. For ease of notations in this paper, we use $G$ to denote this group. Such a definition imparts two important structures on this space, (a) a group structure and (b) a $C^{\infty}$ differentiable structure. This means, we can: (a) construct diffeomorphisms by integrating velocity fields, and (b) combine diffeomorphisms using compositions. This enables us to generate large deformations while maintaining the diffeomorphic property. The smooth differentiable structure on the group of diffeomorphisms makes it a Lie group. A Lie group is a group that is also a smooth manifold. Some of the standard texts to review Lie groups include those by Chevalley (1999) and Adams (1969). In the next section, we discuss the Riemannian structure of the group of diffeomorphisms.

### 3.1.1. Lie algebra and Riemannian structure

The tangent space at identity also called the Lie algebra, $\mathfrak{g}$, of diffeomorphisms, consists of all vector fields with finite norm. Its dual space, $\mathfrak{g}^{*}$, consists of vector-valued distributions over $\Omega$. The velocity, $v \in \mathfrak{g}$, maps to its dual deformation momenta, $m \in \mathfrak{g}^{*}$, via the operator, $L$, such that, $m=L v$ and $v=K m$. The choice of a self-adjoint differential operator, $L$, determines the right-invariant Riemannian structure on the collection of velocity fields with the norm defined as, $\|v\|_{\mathfrak{g}}^{2}=\int_{\Omega}(L v(x), v(x)) d x<\infty$. The diffeomorphisms corresponding to the flows of these velocity field with finite norm constitute a subgroup of $\operatorname{Diff}(\Omega)$. The operator, $K: \mathfrak{g}^{*} \rightarrow \mathfrak{g}$, denotes the inverse of $L$. The Jacobi-Lie bracket of the Lie algebra, which expresses the commutativity of vector fields in $\mathfrak{g}$ can also be written in terms of the adjoint representation operator $\mathrm{ad}_{v} w$ of the Lie algebra. In particular, the bracket takes the form,
$[v, w]=-\mathrm{ad}_{v} w=D v w-D w v$.
The $\operatorname{ad}_{v}$ operator and its conjugate, $\mathrm{ad}_{v}^{*}$, are fundamental to the group. For example, $\mathrm{ad}_{v}^{*}$ determines the geodesic evolution of curves in $G$ via the Euler Poincaré (EPDiff) evolution equation for geodesics (Younes et al., 2009), obtained using the right invariant metric to give the evolution in the Lie algebra such that,
$\dot{m}+\mathrm{ad}_{v}^{*} m=0$.
We provide more details on the derivation of the conjugate of the ad operator in Appendix A.

The operators for the adjoint representation of a group and its Lie algebra are fundamental to the transport of relational structures such as velocities and their duals. Some of the classic texts for a thorough understanding of representation theory include those by Curtis and Reiner (1962) and Fulton and Harris (1991). For a thorough review of the Riemannian structure on the group of diffeomorphisms,

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please refer to Younes (2010), Arnol'd (1966) and Younes et al. (2009). We present the formulation of variational cubic curves in diffeomorphisms in the next section.

### 3.2. Variational formulation

Let us first introduce curves of minimal acceleration on a general Riemannian manifold (Camarinha et al., 1995; Crouch \& Leite, 1995; Noakes et al., 1989) which are needed for the image case. As a boundary value problem, Riemannian cubic splines are defined as curves that minimize the following energy $E(g)=\frac{1}{2} \int_{0}^{1}\left\|\nabla_{\dot{g}} \dot{g}\right\|_{T_{g} G}^{2} d t$ subject to boundary constraints $g(0)=g_{0}, \dot{g}(0)=v_{0}$ and $g(1)=g_{1}, \dot{g}(1)=v_{1}$. Here, $\nabla$ denotes the Levi-Civita connection associated with the Riemannian metric denoted by $\|\cdot\|_{T_{g} G}$. The quantity, $\nabla_{\dot{g}} \dot{g}$, is the generalization of the idea of acceleration to Riemannian manifolds. Note that the associated Euler-Lagrange equation is in the form
$\frac{D^{3}}{D t^{3}} \dot{x}-R\left(\dot{x}, \frac{D}{D t} \dot{x}\right) \dot{x}=0$,
which involves the curvature tensor $R$ associated with the metric. Another way to define the spline is by defining a time-dependent control that forces the curve $g(t)$ to deviate from being a geodesic (Trouvé \& Vialard, 2012). Such a control or a forcing variable, $u(t)$ is then integrated using the formula $\nabla_{\dot{g}(t)} \dot{g}(t)=u(t)$. Notice, $u(t)=0$ implies that $g(t)$ is a geodesic.

Taking this idea forward to the group of diffeomorphisms, $G$, we propose to include a time-dependent forcing term that describes how much the 'geodesic requirement' deviates. Thus, we define the control directly on the known momenta EPDiff evolution equation for geodesics (Younes et al., 2009), obtained using the right invariant metric to give the evolution in the Lie algebra (in fact, the tangent space at the identity deformation), $\mathfrak{g}$, as $\dot{m}+\mathrm{ad}_{v}^{*} m=0$. As mentioned before, the operator ad* is the adjoint of the Jacobi-Lie bracket (Bruveris et al., 2011; Younes et al., 2009). After adding this control, the dynamics take the form, $\dot{m}+\mathrm{ad}_{v}^{*} m=u$, where $u \in \mathfrak{g}^{*}$. Thus we allow the geodesic to deviate from satisfying the EPDiff constraints and constrain it to minimize an energy of the form, $E=\frac{1}{2} \int_{0}^{1}\|u(t)\|_{\mathfrak{g}^{*}}^{2} d t$. It is important to note that such a formulation will avoid direct computation of curvature. We bypass it when we control the EPDiff in $\mathfrak{g}$ instead of controlling $\nabla_{\dot{g}(t)} \dot{g}(t)$ in $T_{g(t)} G$.

The associated constrained energy minimization problem for splines is then,
$\underset{u}{\operatorname{minimize}} \frac{1}{2} \int_{0}^{1}\|u(t)\|_{\mathfrak{g}^{*}}^{2} d t$
$\dot{m}(t)-u(t)+\mathrm{ad}_{K m(t)}^{*} m(t)=0, m(0)=m_{0}$
$\dot{h}(t)+(D h) K m=0, h(0)=i d$
Here we use, $h(t)=g^{-1}(t)$ for ease of notations and id denotes the identity map. Notice that this formulation in diffeomorphisms is analogous to the Euclidean case in the sense that the Euclidean states ( $x_{1}, x_{2}$, $\left.x_{3}\right) \mapsto(h, m, u)$ in diffeomorphisms. The constraint in (5) is the control and (6) is a deformation advection constraint, which describes how the inverse of the diffeomorphism evolves over time. This equation will also play a central role for the matching terms as it allows to match for example, images, landmarks, surfaces, etc.

Similar to the Euclidean case, the Euler-Lagrange equations for the above optimization problem give an adjoint system that explains the evolution of $u$, such that,
$\dot{u}-(D h)^{\top} p-K^{-1} \operatorname{ad}_{K m} K u+\operatorname{ad}_{K u}^{*} m=0$,
$\dot{p}+\nabla \cdot(p \otimes K m)=0$.
Here, $p$ is the adjoint variable corresponding to the deformation evolution constraint on $g^{-1}$ in Eq. (6). The details of the derivation of the
above Euler-Lagrange equations are presented in Appendix B. Note however that the existence of a minimizer to this variational problem is still an open problem whereas the shooting splines solutions introduced in the next section are well-defined.

### 3.3. From relaxation to shooting.

Notice that the above discussion is analogous to the discussion of the relaxation formulation for the Euclidean case. We now convert the adjoint state, $p(t)$ to a primal state to form a forward shooting system. Analogous to the Euclidean case, this increases the order of the system by one. The shooting system for acceleration controlled motion is,

$$
\left.\begin{array}{rl}
\dot{h}+(D h) K m & =0 \\
\dot{m}-u+\mathrm{ad}_{K m}^{*} m & =0 \\
\dot{u}-(D h)^{\top} p-K^{-1} \mathrm{ad}_{K m} K u+\mathrm{ad}_{K u}^{*} m & =0  \tag{9}\\
\dot{p}+\nabla \cdot(p \otimes K m) & =0
\end{array}\right\}
$$

The image evolves (equivalently advects) as per the group action of $g$ on the initial image $I^{0}$, i.e., $I(t)=I^{0} \circ g^{-1}(t)=I^{0} \circ h(t)$. Here, the vector quantity, $p$ is analogous to $x_{4}$.

### 3.4. Shooting-splines with data-independent controls for regression

Similar to the data configuration in the Euclidean example, in the context of regression, let, $J_{i}$, for $i=1 \ldots N$, denote $N$ measured images at timepoints, $t_{i} \in(0,1)$. The goal now is to define finite and relatively fewer points than the number of measurements in the interval, $(0,1)$ where $p$ is allowed to jump. In other words, $p$ does not jump at every measurement but instead, is allowed to be free at predefined timepoints that are decided independently of the data. Thus, we construct a curve $g(t)$, similar to the Euclidean case, in $G$, along the time, $t$, such that it guarantees, (a) $p$ may jump at predefined time-points only, (b) $u$ is $C^{0}$-continuous, (c) $m$ is $C^{1}$-continuous, and (d) $g$ is $C^{2}$-continuous.

The unconstrained Lagrangian for spline regression for a fixed initial image $I^{0}$ takes the form:

$$
\begin{align*}
& E\left(h, m, u, p, p^{t_{c}}, \lambda_{h}, \lambda_{m}, \lambda_{u}, \lambda_{p}\right)=  \tag{10}\\
& \frac{1}{2 \sigma_{I}^{2}} \sum_{c=1}^{C+1} \sum_{i \in \mathcal{I}_{c}} \mathrm{~d}^{2}\left(I^{0} \circ h\left(t_{i}\right), J_{i}\right)+\frac{1}{2 \sigma_{u}^{2}} \int_{0}^{1}\langle u, K u\rangle_{L^{2}} d t \\
& \quad+\sum_{c=1}^{C+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{h c}, \dot{h}_{c}+\left(D h_{c}\right) K m_{c}\right\rangle_{L^{2}} d t \\
& \quad+\sum_{c=1}^{C+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{m c}, \dot{m}_{c}-u_{c}+\mathrm{ad}_{K m_{c}}^{*} m_{c}\right\rangle_{L^{2}} d t \\
& \quad+\sum_{c=1}^{C+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{u c}, \dot{u}_{c}-\left(D h_{c}\right)^{\top} p_{c}-K^{-1} \mathrm{ad}_{K m_{c}} K u_{c}+\mathrm{ad}_{K u_{c}}^{*} m_{c}\right\rangle_{L^{2}} d t \\
& \quad+\sum_{c=1}^{C+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{p c}, \dot{p}_{c}+\nabla \cdot\left(p_{c} \otimes K m_{c}\right)\right\rangle_{L^{2}} d t, \text { and }
\end{align*}
$$

subject to continuity of $h, m$ and $u$ at $C$ joins.
Note that we use subscript $c$ to denote that the variables, $h_{c}, m_{c}, u_{c}$ and $p_{c}$, and their corresponding adjoints are defined in their respective intervals $\mathcal{I}_{c}$ and they stitch together to give $h, m, u$ and $p$, defined in $(0,1)$. Also notice the second term is the relaxation energy term, i.e., $\|u\|_{\mathfrak{g}^{*}}^{2}=\langle u, K u\rangle_{L^{2}}$, which acts as a regularizer on the force along the full path in diffeomorphisms. It is helpful to make a note of the underlying spaces for the variables parametrizing the spline. In particular, $h$ $\in G$, while all the other states are in the coalgebra, i.e., $m, u, p, p^{t_{c}} \in \mathfrak{g}^{*}$.

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### 3.5. Gradients

The optimality conditions on the gradients of the above energy functional show that the adjoint variables, $\lambda_{h}, \lambda_{m}, \lambda_{u}$, are continuous at all $C$ joins. The gradients with respect to the initial conditions are,
$\delta_{m_{1}^{0}} E=-\lambda_{m 1}(0), \delta_{u_{1}^{0}} E=-\lambda_{u 1}(0), \delta_{p_{1}^{0}} E=-\lambda_{p 1}(0)$.
We compute the gradients by integrating the adjoint system of equations within each interval backward in time,

$$
\left.\begin{array}{r}
\dot{\lambda}_{h c}-\nabla \cdot\left(p_{c} \otimes \lambda_{u c}-\lambda_{h c} \otimes K m_{c}\right)=0 \\
\dot{\lambda}_{m c}-\operatorname{ad}_{K m_{c}} \lambda_{m c}-\operatorname{ad}_{K u_{c}} \lambda_{u c} \\
+K \operatorname{ad}_{\lambda_{m c}}^{*} m_{c}-K \mathrm{ad}_{K u_{c}}^{*} K^{-1} \lambda_{u c} \\
+K\left(\left(D \lambda_{p c}\right)^{\top} p_{c}\right)-K\left(\left(D h_{c}\right)^{\top} \lambda_{h c}\right)=0  \tag{11}\\
\dot{\lambda}_{u c}-\frac{1}{\sigma_{u}^{2}} K u_{c}+\lambda_{m c}+K \operatorname{ad}_{K m_{c}}^{*} K^{-1} \lambda_{u c} \\
+K \operatorname{ad}_{\lambda_{u c}}^{*} m_{c}=0 \\
\dot{\lambda}_{p c}+\left(D h_{c}\right) \lambda_{u c}+\left(D \lambda_{p c}\right) K m_{c}=0
\end{array}\right\}
$$

The details of the derivation of the above Euler-Lagrange equations are presented in Appendix C.

All variables start from zero as their initial conditions for this backward integration. Similar to the Euclidean case, we add jumps in $\lambda_{h}$ as,
$\lambda_{h c}\left(t_{i}^{+}\right)-\lambda_{h c}\left(t_{i}^{-}\right)=\frac{1}{\sigma_{I}^{2}}\left(I^{0} \circ h\left(t_{i}\right)-J_{i}\right) \nabla I^{t_{i}}$,
at measurements, $t=t_{i}$ if we use a sum-of-squared differences similarity measure for $d^{2}(\cdot, \cdot)$ between the measured images $\left\{J_{i}\right\}$ and the estimated images $\left\{I^{0} \circ h\left(t_{i}\right)\right\}$. More general similarity measures could easily be used and would only change these jump conditions. We ensure the continuity of $\lambda_{h c}, \lambda_{m c}$, and $\lambda_{u c}$ at the joins and $\lambda_{p c}$ starts from zero at every join. We use the accumulated $\lambda_{p c+1}$ to update the jerk, $p_{c}\left(t_{k}\right)$, at the control location with,
$\delta_{p_{c+1\left(t_{c}\right)}\left(t_{c}\right)} E=-\lambda_{p c+1}\left(t_{c}^{+}\right)$.
Note this is the 'data independent' control that we motivated our formulation with. This determines the initial condition of the forward system for each interval and needs to be estimated numerically. Also note that other regularizers can be added on the initial momenta, $m_{1}^{0}$, initial jerk, $p_{1}^{0}$, and jerks at controls, $p_{c+1}^{t_{c}}$, by restricting their Sobolev norms. In this case, the gradient includes additional terms of the form, $K m_{1}^{0}, K p_{1}^{0}$, and $K p_{c+1}^{t_{c}}$, respectively. The estimate for the force term, $u_{0}$, does not need to be regularized since minimizing the norm on $u(t)$ along the path itself acts as a regularizer. Finally, all the above gradients generalize trivially in case we have data measurements at the end points or at control locations. This results in additional jumps (limit from the right) in the adjoint variable at these measurement locations during the backward integration.

## 4. Results

We evaluate our proposed model using synthetic data and two real time-sequence imaging data sets. One of the real imaging data examples is from cellular imaging of snapshots acquired for a deforming cell imaged using atomic force microscopy. The other is from the Sunnybrook cardiac MR database (Radau et al., 2009). In all our experiments, we fix the initial image, $I(0)$, and estimate initial states, $m(0)$, $u(0), p(0)$ and $p\left(t_{c}\right)$ at control locations that completely determine the spline curve $g(t)$ from $t=0$ to $t=1$.

Implementation details and discretization schemes. The experiments constitute of estimating the parameters of the best fitting regression curves in diffeomorphisms for a sequence of images, as well as of generating or synthesizing the new curves based on the given parameters
independent of any image data. The primary component for implementing the cubic and spline curve estimation involves the forward integration of the set of ODEs in Eq. (9), representing the forward curve evolution, and the backward integration of the set of ODEs in Eq. (11), representing the adjoint evolution. The adjoint states in the adjoint system, in turn, provide the gradients for updating the parameters of the curves during the optimization. Note that synthesizing curves given the parameters only involves integrating the forward system in Eq. (9). The terms in all of these ODEs including the terms with 'ad' operators, involve computing finite difference approximations of the pixel/voxel-level spatial gradients, divergences, and their combinations.

We use a fourth order Runge-Kutta method with 40 timesteps to integrate the primal states forward and to integrate the corresponding adjoint states backwards. We use finite difference approximations with central differences for all pixel/voxel-level gradient, Jacobians and divergence computations. Also, note that for computing the adjoint of the Jacobi-Lie bracket, $\mathrm{ad}_{v}^{*} m=(D v)^{T} m+\nabla \cdot(m \otimes K m)$, the second term is essentially the divergence of the tensor field. The tensor divergence of the field $S$ is expressed using Einstein summation as,
$\nabla \cdot S=\frac{\partial S_{k i}}{\partial x_{i}} e_{k}$.
In the case where the tensor field is represented as a pixel/voxel-wise Kronecker product of 3D vector fields, $u$ and $v$, the tensor divergence takes the form,
$\nabla \cdot(u \otimes v)=\left(\begin{array}{l}\frac{\partial u_{1} v_{1}}{\partial x}+\frac{\partial u_{1} v_{2}}{\partial y}+\frac{\partial u_{1} v_{3}}{\partial z} \\ \frac{\partial u_{2} v_{1}}{\partial x}+\frac{\partial u_{2} v_{2}}{\partial y}+\frac{\partial u_{2} v_{3}}{\partial z} \\ \frac{\partial u_{3} v_{1}}{\partial x}+\frac{\partial u_{3} v_{2}}{\partial y}+\frac{\partial u_{3} v_{3}}{\partial z}\end{array}\right)$.
Again, we use finite difference approximations with central differences for evaluating the above spatial partial derivatives.

We use a line search with gradient descent to estimate the optimal initial states of spline curves and the controls and the kernel, $K$, corresponds to the invertible and self-adjoint Sobolev operator, $L=$ $-j \nabla^{2}-k \nabla(\nabla \cdot)+l$, with $j=0.2, k=0.2$, and $l=0.001$.

A typical runtime for executing a full line search gradient descent optimization algorithm on a sequence of $102 \mathrm{D} 84 \times 84$ sized images is about 20 min for 100 iterations on the Quadro K200 NVIDIA GPU. The CPU-GPU implementations of the shooting splines for 2D and 3D image sequences can be obtained from http://git@bitbucket.org:nikhilsingh/diffeosplines.git.

In our validations, we first experimentally demonstrate that the shooting equations for evolving cubic curves in diffeomorphisms as per the set of ODEs in Eq. (9) are indeed analogous to the classical system of shooting cubic curves in Euclidean as given in Eq. (2) and to the cubic equation, $y=a x^{3}+b x^{2}+c x+d$. We discuss these results in Sec. 4.1. Next, we present our experiments for regression with splines using synthetic data, the deforming cell imaging data and the cardiac imaging data in Sec. 4.2.

### 4.1. Assessment of shooting higher order curves

We first study the interpretation of the new states proposed in this paper: the force denoted by $u(t)$ and the jerk denoted by $p(t)$ for the evolution of curves in diffeomorphisms. It is informative to study the simplest case of regression first: the image matching problem between a set of two images, such that it solves the variational problem in Eq. (10) in the absence of any control locations, i.e., $C=0$. Since, there are only two points, the energy minimizing curve is a geodesic path but one that allows changes in velocity and in acceleration. The

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Fig. 4. Higher order matching of two images. (a) Top row is the momenta only matching which is equivalent to the usual geodesic matching of images such that $u(t)=0$ and $p(t)=$ 0 along the matching path in diffeomorphisms. This matching only estimates initial momenta, $m(0)$. (b) Middle row is for force only matching to estimate $u(0)$ such that $p(t)=0$ along the path but $m(t)$ accelerates from zero initial condition, i.e., $m(0)=0$. This matching only estimates initial force, $u(0)$. (c) Bottom row corresponds to jerk only matching such that both $m(t)$ and $u(t)$ accelerate from zero initial conditions, i.e., $m(0)=u(0)=0$. This matching only estimates initial jerk, $p(0)$. The length of arrows is proportional to their scales. On the right is the convergence of gradient descent with line search for the three optimization problems.
fourth derivative, jerk, however must conserve mass during its transport. Also note that if we only constrain the jerk state to be zero at all times, such that $p(t)=0$, the diffeomorphism evolves in a "quadratic" form. Along with the jerk, if we also constrain the force state to be zero at all times, such that $u(t)=0$, the diffeomorphism evolves in the standard geodesic form.

To investigate this, we decompose our analysis into three simpler experiments. We first solve the problem of matching two images by constraining the force and the jerk states to be zero at all times and only estimate the initial momenta that describe the geodesic. We call this the momenta only matching (Fig. 4, top row). Images on the left are the initial states: image, momenta, force and jerk at $t=0$, and on the right are the final states: image, momenta, force and jerk at $t=1$. This is simply solving the classical image matching problem using the geodesic shooting equations. In this case, the momenta evolution satisfies the standard EPDiff evolution. In the view of classical mechanics and particle motion, it is also analogous to describing the motion of a particle with constant velocity under the absence of any external force. Next, we solve the problem of matching the same images by constraining the starting momenta to be zero at $t=0$, and the jerk states to be zero at all times, and estimate only the initial force. We call this the force only matching (Fig. 4, middle row). This is analogous to the motion of a stationary particle under constant force such that it starts from zero velocity and then constantly accelerates. Finally, we solve the problem of matching the same two images by constraining the starting momenta and force to be zero at $t=0$, and estimate only the initial jerk. We call this the jerk only matching (Fig. 4, bottom row). This is analogous to describing the motion of a stationary particle with a continuous impulse such that it starts moving from zero velocity, and zero acceleration and then its acceleration increases constantly and its velocity increases in second order. The simple gradient descent with line search optimization converges for these three experiments. As expected, the momenta at the end point of the matching path for the jerk only matching are larger than that observed at the end point for force only matching. Also, the final force state for jerk only matching ends up being larger in magnitude than the force state for force only matching. Note that for all three curves the start point (the identity deformation) and the end points (the best matching deformation) are identical. The matching only differs in the order of motion along the path the curve traces in diffeomorphisms.

To further understand the Euclidean analogy of our proposed shooting equations in Eq. (9) we combine the three estimates and observe the resulting evolution. In particular, the above three experiments result in the three coefficients that follow similar scaling rules as the standard cubic curves, $y=a x^{3}+b x^{2}+c x+d$, where $a$ is equivalent to $p(0)$ estimated for jerk only matching, $b$ to $u(0)$ estimated for force only matching, and $c$ to $m(0)$ obtained for momenta only matching. We can now conveniently synthesize different parametric curves using the scaling of these coefficients. In Fig. 5, we demonstrate the quadratic and the cubic polynomial curves in diffeomorphisms synthesized using these estimates obtained from matching.

To simulate a curve similar to the Euclidean quadratic, $y=x^{2}-$ $x$, we integrate Eq. (9) starting from the initial momenta of $m(0)$ and initial force of $-u(0)$. We observe that the diffeomorphic path traces a quadratic path which is analogous to the motion of a particle under constant force. This is similar to the motion of a particle with a given initial velocity at $t=0$ but opposite force such that the particle decelerates initially, comes to a rest state exactly at $t=0.5$ and then accelerates to return back to the exact initial position at $t=1.0$ (Fig. 5(a)). The shooting equations are accurate such that the diffeomorphisms end at the identity transformation at $t=0$. Another way to visualize this path in diffeomorphisms is to observe the motion of a pixel at the boundary of the image as it deforms from $t=0$ to $t=1$. For this we display stacked-up 1D cross sections (middle row of the image). This forms a 2D matrix displayed as a picture in Fig. 5(c), such that the rows are stacked up in increasing order of time from bottom to top. We notice that the pixels trace a quadratic curve as the image deforms along this path.

To demonstrate a cubic-like behavior, we scale the coefficients to generate a Bernoulli polynomial of degree 3 that takes the form, $y=$ $x^{3}-\frac{3}{2} x^{2}+\frac{1}{2} x$. For this, we integrate Eq. (9) starting with scaled initial conditions in the same proportion as the Bernoulli coefficients and use the initial momenta of $m(0)$, initial force of $-3 u(0)$ and the initial jerk of $2 p(0)$. We observe that the diffeomorphic path traces a cubic path such that the shape first compresses in one direction and then expands and reaches back to a state that it started with and continues to expand until it finally shrinks back (Fig. 5(b)). This is analogous to the behavior a Euclidean Bernoulli polynomial of degree 3 follows with the exact same coefficients. Also, similar to the visualization of the quadratic curve, the cross sectional boundary pixel visualization

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Fig. 5. Quadratic and cubic polynomial paths in diffeomorphisms. Row (a) corresponds to the diffeomorphic path for quadratic evolution obtained by shooting along $m(0)$ and $-u(0)$. This is analogous to the motion of a particle with finite initial velocity under constant decelerating force. Row (b) corresponds to the cubic evolution obtained by shooting along coefficients scaled as per the Bernoulli coefficients, i.e., $m(0),-3 u(0)$ and $2 p(0)$. Bottom row, (c) and (d) are the cross-section visualizations of the generated images along these paths. The trajectory of boundary points follows quadratic and cubic paths, respectively. Note that the base coefficients, $m(0), u(0)$ and $p(0)$ that were scaled, are the initial conditions estimated by solving the image registration problem with momenta only, force only and jerk only constraints, respectively, as per Fig. 4.
of this curve results in a Bernoulli path that resembles a Bernoulli cubic (Fig. 5(d)).

### 4.1.1. Quadratic and cubic regression

Next, to assess the strength of regression using these parametric shooting equations, we generate ( $N=9$ ) sampled shapes to simulate non-monotonic quadratic and cubic like dynamics from $t=0$ to $t=1$ (Fig. 6(a) and (c), respectively) using the shooting methods described above. The corresponding quadratic and cubic regression fit are shown in Fig. 6(b) and (d), respectively.

A note on initialization. A good initialization is necessary since the variational problem is non-convex. A possible strategy for the initialization is to first compute the initial momenta, $\tilde{m}$, for matching the first image (source image) with the next image in sequence that looks to be the most deformed image (target image) relative to the initial image. In the case of quadratic data in row (a), a good candidate for the target image could be the one at $t=0.5$ while for the cubic data in row (c), a possible candidate for the target image could be the one at $t=0.25$. A good initialization for the quadratic regression gradient descent optimization problem could then use this momenta direction estimated only from the data to initialize both the initial quadratic states, for example, using approximately scaled $\tilde{m}$ : $m^{k}(0)=2 \tilde{m}$ and $u^{k}(0)=-4 \tilde{m}$ at first iteration for $k=1$. Similarly, for the cubic regression variational problem, such an initialization could be used to also initialize all the initial states of momenta, force and jerk for the first iteration at $k=1$ of the gradient descent, for example, using approximately scaled: $m^{k}(0)=2 \tilde{m}, u^{k}(0)=-12 \tilde{m}$ and $p^{k}(0)=24 \tilde{m}$, where $\tilde{m}$ denotes the momenta corresponding to the geodesic matching problem of matching initial image with the image at $t=0.25$. Note these factors correspond to quadratic and Bernoulli cubics respectively along with additional scalings to compensate for the shorter length of the path for which $\tilde{m}$ is computed.

We emphasize that the above strategies use only the data to decide initial conditions of a curve that could be close to the possible minimizing least square polynomial curve. This provides a practical way to find good initialization for the fitting problem for any given
data set. We notice that using such initializations the optimization converges within 100 iterations for both the quadratic and the cubic regression variational problems (Fig. 6(f) and (h)).

For the quadratic regression, we notice that the estimated fit captures the trend and results in a smoothly shrinking followed by expanding grids along the regression path to closely match the data. The boundary pixel also traces a quadratic curve when the image deforms (Fig. 6(e)). Similarly, for the cubic regression, the estimated fit captures the two inflections of the motion to best fit the data. The boundary pixel also traces a quadratic curve when the image deforms (Fig. 6(e)).

### 4.2. Assessment of spline regression

In this section, we investigate the performance of spline regression on synthetic and real data. For the synthetic data, we generate the data such that a cubic-like dynamic alone is not sufficient to explain the trends in shape changes and therefore necessitates adding a control for the spline fit. For the real data, we perform spline regression experiments using cell and cardiac images.

### 4.2.1. Synthetic data

To assess the strength of spline regression for non-geodesic image data, we create a synthetic sequence of $N=13$ shapes to simulate nonmonotonic changes with more than two inflection points from $t=0$ to $t=1$ (Fig. 7(a)). The synthetic shape first shrinks and then expands till $t=0.5$, and then again shrinks and finally expands back again till it reaches the end point, at $t=1.0$. The last image in the sequence in Fig. 7(a), shows another visualization in the form of stacked-up 1D cross sections (middle row of the input images). This forms a 2D matrix displayed as a picture, such that the rows are stacked up in increasing order of time from bottom to top with blank rows for the times for which there is no data. Using such data, we attempt to simulate dynamics such that a cubic alone would not be sufficient to trace through the inflection. For optimization, we follow a similar strategy we discussed in Section 4.1.1 for cubic regression to initialize

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Fig. 6. Regression of quadratic and cubic like data. Top two rows (a) and (b) detail the quadratic like sparsely sampled data and the corresponding quadratic regression fit, respectively, and (e) and (f) show the cross-sectional view of movement of a boundary point along the quadratic fit and the convergence of the optimization, respectively. Similarly, rows (c) and (d) detail the cubic-like sparsely sampled data and the corresponding cubic regression fit, respectively, and (g) and (h) show the cross-sectional view of movement of a boundary point along the cubic fit and its convergence, respectively.

Table 1
Total squared error of fit.

|  | Spline fit (one control) | Spline fit (no control) | Geodesic fit |
| :--- | :---: | :---: | :---: |
| Synthetic | 40.50 | 139.77 | 162.39 |
| Cell data | 70.68 | 71.36 | 307.18 |
| Cardiac data | 962.75 | 975.15 | 1440.58 |

momenta, force and jerk at $t=0$ but just switch the sign on the jerk state, $p$ at the control location to add another inflection.

We report a quantitative comparison of the three fits in the first row of Table 1. The reported error of fit corresponds to $L^{2}$ image residual as per the data-likelihood in Eq. (10). We observe that adding a single spline control at the mid point results in the best fit that summarizes the smooth dynamics of change (Fig. 7(b)). The estimated diffeomorphism successfully captures the two cycles of trends in the shrinking shape, followed by expansion. Without adding any control, the resulting spline trend, even though a cubic, fails to capture the dynamics and fails to recover the inflection points in the rate of shape change (Fig. 7(c)). Finally, being the most inflexible, geodesic regression performs worst, and barely captures any real spatio-temporal trend (Fig. 7(d)). The visualization of the motion of boundary pixels also confirms (last column) the flexible diffeomorphism fit obtained for the spline regression with one control. For further details on the analysis of the regression fit, please see the visualizations in the supplementary material that include the gif video of the fit and the mosaic overlay of the original data with the regressed trend.

### 4.2.2. Deforming cell data

The cell time-sequence data corresponds to 11 snapshots at equal intervals a deforming cell imaged using an atomic force microscope. We preprocessed the images using total variation denoising (Chambolle, 2004) to preserve edges (Fig. 10). Note that the original images are noisy with a grainy texture while the denoised image effectively retains the regions of high gradients, which are useful for driving the spline regression. The images depict a trend in which the shape of the cell deforms such that its left boundary first bends inward and then resumes back to its original shape (Fig. 8(a)).

A visual assessment of the cell images suggests that this data should have an inflection in the dynamics of shape changes. The regression fit using one control and no control results in a very similar fit (Fig. 8(b) and (c)). However, the geodesic fit only results in a monotonous compression of the cell and fails to capture the expansion in the last half of the dynamics (Fig. 8(d). We also notice in Table 1 that the spline regressions with one control and without a control result in comparable $L^{2}$ error of fit as per the data likelihood. The geodesic fit, however, clearly performs worst in terms of the residual error of fit. We present more details on the analysis of the regression fit in the visualizations in the supplementary material that include the gif video of the fit and the mosaic overlay of the original data with the regressed trend.

### 4.2.3. Cardiac data

The cardiac time-sequence data corresponds to 20 snapshots at equal intervals of the beating heart of a normal individual with age $=$ 63 years (Subject Id: SCD0003701). We cropped all the axial images to

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(a) Original data

(b) Spline fit (control at $t=0.5$ )

(c) Spline fit (no control)

(d) Geodesic fit


Fig. 7. Comparison of regression models on the synthetic data. (a) The original synthetic shapes that go through non-monotonic deformations simulating three inflection points. The last image in this sequence visualizes the pixel at the boundary, in the middle 1D horizontal slice of each of these 2D input image, such that the black region represents no data. (b) The spline fit using a control at location $t=0.5$ overlaid with the corresponding deformed grid. The last image in the sequence visualizes the motion of a pixel at the boundary, in the middle 1D horizontal slice of the deforming 2D image, as we move along the regression fit in diffeomorphisms. (c) and (d) The spline and the geodesic fit, respectively.
a common rectangular region around the heart followed by histogram matching to align intensities of all the timepoints to the image at $t=0$. Fig. 9 shows the original scans (first row) and the result of regression models (second to fourth row). We only display half of the timepoints of the ones actually used to fit the model. Similar to the synthetic data, we observe that the original data exhibits a non-geodesic and non-monotonic trend in changing shape of the beating heart. The comparison in terms of the error of fit for all models suggests that both spline curves perform better than the geodesic. Although visually the fits look similar for splines, we obtained a marginal improvement of fit for the spline curve with single control when compared to the
spline curve without any control (Table 1). The geodesic fit again performs worst out of the three models. The dynamics of the beating heart for these models are best seen in the multimedia file in the supplementary material and the mosaic overlay image of the original data with the regressed trend.

## 5. Discussion

In this article, we developed a theory for higher order curves that generalizes the notion of parametric curves such as the quadratic, the cubic and the piecewise cubics to the manifold of
(a) Original data

(b) Spline fit (control at $t=0.5$ )

(c) Spline fit (no control)

(d) Geodesic fit


Fig. 8. Comparison of regression models on the deforming cell data. (a) The original cell images that shrink and then expand back. (b) The spline fit using a control at location $t=$ 0.5 overlaid with the corresponding deformed grid. (c) and (d) The spline with no control and the geodesic fit, respectively.
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(a) Original data

(b) Spline fit (control at $t=0.5$ )

(c) Spline fit (no control)

(d) Geodesic fit


Fig. 9. Comparison of regression models on the cardiac motion data. (a) The original cardiac snapshots. (b) The spline fit using a control at location $t=0.5$ overlaid with the corresponding deformed grid. (c) and (d) The spline with no control and the geodesic fit, respectively.
diffeomorphisms. We provided a principled way to define curves with nonzero acceleration and nonzero jerk, which is the natural next step of extension to geodesic-based methods developed during the last decade for computational anatomy in the large deformation diffeomorphic image analysis framework. We took a variational approach that is governed by an underlying energy formulation, which respects the nonflat geometry of diffeomorphisms. Such an approach of minimal energy curve estimation also provides a physical analogy with particle motion under a varying force field.

As a consequence, the initial conditions of our variational quadratics and cubics are interpretable similar to the initial conditions of their corresponding Euclidean parametric counterparts. To validate this, we demonstrated that evolving the curves according to scaled initial conditions also results in the same behavior as a scalar Euclidean parametric curve would exhibit under these scalings. We tested this for different scalings and presented the results for quadratic parabolic scalings and for Bernoulli cubic scalings.

Our proposed system of evolution equations for higher order curves can be used for regression. The benefit of using these forward


Fig. 10. An example of the cell image data before and after the total variation denoising.
shooting equations in an optimal control setting is that the solution to the resulting regression problem is given only in terms of a few initial conditions. We emphasize that in all our experiments, the full diffeomorphic paths and the evolution of all states along the estimated curves are completely parameterized by very few parameters (four for spline fits with one control and three for spline fits without any control) that are independent of the study size.

### 5.1. Open questions and extensions

Regression models are expected to fit better with an increasing number of control points. This necessitates model selection procedures. In other words, a criterion to assess the balance of increasing model complexity to obtain better data fit needs to be developed. Euclidean methods of model selection such as the Akaike information criterion (AIC) and the Bayesian information criteria (BIC), both require the notion of distribution and hence are difficult to generalize to diffeomorphisms. Nonparametric approaches such as permutation tests may provide a possible way to define and approximate null distributions empirically and to test the significance of models for a given dataset. Alternatively, cross-validation could be used for model selection.

Due to the limitations of our current optimization method, i.e., gradient descent, our spline estimation experiments used at most one control point only. A possible future work would be to explore better optimization strategies and to develop second order methods utilizing limited memory for the optimization of splines (Byrd et al., 1995). We expect improved convergence for example by using a quasi-Newton methods such as IBFGS.

Note that adding an extra control point may not always result in a noticeable improvement for practical scenarios. The need for control points will be data-set dependent. For example, adding a control point generally shows clear improvements for the cases when the data exhibits more than one inflection point. However, in our experiments both the real datasets exhibit trends that show only one directional change in velocity, e.g., the ventricles in the heart expand and then shrink for only one cycle. Similarly, the cell exhibits a shrinkage followed by an expansion back to its original position. However, our synthetic example clearly demonstrates the benefit of adding a control point for a dataset that has more than one inflection point. Hence, we expect control points also to be useful for real data for sufficiently complex deformations.

We developed the regression problem for a fixed initial image. Adding template estimation would add another parameter to the estimation problem. With better optimization strategies, it should also be possible to develop an alternate optimization algorithm for template estimation (Singh et al., 2013b).

The position of control points is also a modeling choice. The uniform placement of control points on the axis of the regressor variable is convenient and facilitates model comparisons and interpretability in medical imaging population studies. However, it remains an open problem to also optimize for the locations of control points. The possible challenges will include investigating the differentiability of the
energy functional in Eq. (10) with respect to the location of the jump in the jerk state.

Another aspect could be investigating possibilities of combining our model on diffeomorphisms with the higher order models on shapes (Gay-Balmaz et al., 2012; Trouvé \& Vialard, 2012).

One of the most critical contributions in this research is the ability to trace a path in diffeomorphisms with nonzero acceleration. This could be immensely useful for medical studies of growth or decline where the rate of change gets affected and the emphasis is on accelerated tissue growth or decline. One future possible application for the quadratic models could be to study differences in aging of individuals with or without dementia and investigating the ages and local region in the brain exhibiting the most accelerated atrophy. This would add second order information on tissue atrophy to the information currently being obtained using contemporary first order geodesic regression methods and deformation based morphometry analysis.

Another use of higher order models is for the recently proposed longitudinal models for diffeomorphisms such as hierarchical geodesic models (HGM) (Singh et al., 2013a). Even though for brain studies, a geodesic-like trend is expected to be a good approximation of changes in the brain for a single individual when the measurements are taken within a span of five years, the geodesic assumption on the average group trend of the entire population data for staggered designs may not be the best modeling choice. Thus, our proposed higher order curves would provide a better model for the longitudinal summary of a group spanning a wide range of ages from, say, 60-90 years.

Our method of shooting splines in diffeomorphisms lays a foundation to model flexible dynamics of shape changes seen in time series of medical images, and also opens the possibility to model periodic data by adding periodicity constraints.

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## Appendix A. Conjugate of the Lie algebra adjoint representation

We denote the Jacobian operator by $D(\cdot)$ and the pixel/voxel-wise Kronecker product of two vectors by $\otimes$. We use the Riesz representation theorem to first define the dual pairing as a linear operation for integrations on domain $\Omega$ (Rudin, 1986, 1991). For a scalar function, $f \in C_{c}(X)$, let $X$ be a locally compact Hausdorff space. For any positive linear functional, $\psi$ on $C_{c}(X)$, there is a unique Borel regular measure, $\mu$ on $X$, such that,
$\psi(f)=\int_{X} f(x) d \mu(x)$.
We extend it for our case where $f$ is continuous vector valued, say $v$, and not a scalar function. The dual pairing written for the $m(v)$ as $(m, v)$ is written as:
$(m, v)=\int_{\Omega} v(x) d m$.
Notice, $m=u \mu$ where $u$ is a smooth vector field and $\mu$ is the Lebesgue measure on $\Omega$. The above pairing can also be written as:
$(u \mu, v)=\int_{\Omega} v(x) \cdot u(x) d \mu$,
where '. ' is the dot product operation.

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The definition of the conjugate of $\mathrm{ad}_{w}$ is:
$\left(\mathrm{ad}_{w}^{*} m, v\right)=\left(m, \mathrm{ad}_{w} v\right)$.
The right hand side of the above equation takes the form,
$\left(m, \mathrm{ad}_{w} v\right)=\int_{\Omega} \mathrm{ad}_{w} v(x) d m$,

$$
=\int_{\Omega} \operatorname{ad}_{w} v(x) \cdot u(x) d \mu .
$$

The conjugation of the operator, $\mathrm{ad}_{w}$, is computed as follows:

$$
\begin{aligned}
\left(\operatorname{ad}_{w}^{*} m, v\right) & =\int_{\Omega}(D w v-D v w)(x) \cdot u(x) d \mu \\
& =\int_{\Omega}\left((D w)^{T} u\right)(x) \cdot v(x) d \mu \\
& -\int_{\Omega}(D v w)(x) \cdot u(x) d \mu
\end{aligned}
$$

We have isolated, $v$ for the first term. For the second term, we use the relation of the Frobenius norm, where for a matrix $A$, and vectors, $b$ and $c$,
$A b \cdot c=\langle A, c \otimes b\rangle_{\mathcal{F}}$,
where $\mathcal{F}$ represents the Frobenius inner product. Thus,

$$
\begin{aligned}
\left(\operatorname{ad}_{w}^{*} m, v\right) & =\int_{\Omega}\left((D w)^{T} u\right)(x) \cdot v(x) d \mu \\
& -\int_{\Omega}\langle D v(x), u(x) \otimes w(x)\rangle_{\mathcal{F}} d \mu
\end{aligned}
$$

Moreover, since the conjugate of the Jacobian operator, $D$, is the negative of the divergence, we obtain,

$$
\begin{aligned}
\left(\operatorname{ad}_{w}^{*} m, v\right) & =\int_{\Omega}\left((D w)^{T} u\right)(x) \cdot v(x) d \mu \\
& +\int_{\Omega}\langle v(x), \nabla \cdot(u(x) \otimes w(x))\rangle_{\mathcal{F}} d \mu .
\end{aligned}
$$

The above relation implies that as an operator,
$\mathrm{ad}_{w}^{*} \mu=(D w)^{T} \mu+\nabla \cdot(\mu \otimes w)$.
Note that the second term is the pixel/voxel-wise tensor divergence and it can further be split as,
$\nabla \cdot(\mu \otimes w)=D \mu w+\mu \nabla \cdot w$.

## Appendix B. Euler-Lagrange for relaxation problem

We determine the Euler-Lagrange equation in the context of the regression problem. We write the problem of regression for the fixed initial image $I^{0}$ and use $h(t)$ to denote the inverse deformation $g^{-1}(t)$. For readability we drop the argument, $t$, for all the time dependent states, $h(t), m(t)$ and $u(t)$. The constrained energy minimization for the relaxation problem of minimizing the elastic energy for the force controlled curve takes the form,

$$
\begin{aligned}
\tilde{E}\left(h, m, u, \lambda_{h}, \lambda_{m}, \lambda_{u}\right)= & \frac{1}{2} \int_{0}^{1}\langle u, K u\rangle_{L^{2}} d t \\
& +\int_{0}^{1}\left\langle\lambda_{h}, \dot{h}+(D h) K m\right\rangle_{L^{2}} d t \\
& +\int_{0}^{1}\left\langle\lambda_{m}, \dot{m}-u+\mathrm{ad}_{K m}^{*} m\right\rangle_{L^{2}} d t
\end{aligned}
$$

where $K$ correspond to the time independent metric kernels for $u$ and $m$, and $\mathrm{d}(\cdot, \cdot)$ is the metric on images. Note that the metric on images can also be the metric on $G$, here we will keep this simpler, and derive our results for the $L^{2}$ metric.

## B1. Variations

In what follows, unless specified otherwise, all inner products correspond to the $L^{2}$ pairing. Computing variations of $\tilde{E}$ with respect to adjoint variables will give us the dynamic constraints back. Computing variations of $\tilde{E}$ with respect to all state variables gives,

$$
\begin{aligned}
\delta_{h} \tilde{E}= & \left\langle\lambda_{h}(1), \delta h(1)\right\rangle-\left\langle\lambda_{h}(0), \delta h(0)\right\rangle \\
& +\int_{0}^{1}\left\langle-\dot{\lambda}_{h}-\nabla \cdot\left(\lambda_{h} \otimes K m\right), \delta h\right\rangle d t \\
\delta_{m} \tilde{E}= & \left\langle\lambda_{m}(1), \delta m(1)\right\rangle-\left\langle\lambda_{m}(0), \delta m(0)\right\rangle \\
& +\int_{0}^{1}\left\langle-\dot{\lambda}_{m}+K\left((D h)^{\top} \lambda_{h}\right)+\operatorname{ad}_{K m} \lambda_{m}\right. \\
& \left.-K \mathrm{ad}_{\lambda_{m}}^{*} m, \delta m\right\rangle d t \\
\delta_{u} \tilde{E}= & \int_{0}^{1}\left\langle K u-\lambda_{m}, \delta u\right\rangle d t
\end{aligned}
$$

At the optimum the above must vanish. This results in the following adjoint system,

$$
\begin{aligned}
\dot{\lambda}_{h}+\nabla \cdot\left(\lambda_{h} \otimes K m\right) & =0 \\
\dot{\lambda}_{m}-K\left((D h)^{\top} \lambda_{h}\right)-\operatorname{ad}_{K m} \lambda_{m}+K \mathrm{ad}_{\lambda_{m}}^{*} m & =0 \\
K u-\lambda_{m} & =0
\end{aligned}
$$

Notice that, similar to the Euclidean case where the adjoint variable corresponding to the second state, $x_{2}$, turned out to be equal to the third state $x_{3}$, here $\lambda_{m}=K u$.

$$
\dot{\lambda}_{h}+\nabla \cdot\left(\lambda_{h} \otimes K m\right)=0
$$

$K \dot{u}-K\left((D h)^{\top} \lambda_{h}\right)-\operatorname{ad}_{K m} K u+K \operatorname{ad}_{K u}^{*} m=0$
Again, analogous to the Euclidean state, where the adjoint variable corresponding to the first state, $x_{1}$, was renamed as the primal state, $x_{4}$, for the shooting evolution, we rename $\lambda_{h}$ to be $p$ to describe the evolution of $u$ for shooting cubics to give:

$$
\begin{array}{r}
\dot{p}+\nabla \cdot(p \otimes K m)=0 \\
K \dot{u}-K\left((D h)^{\top} p\right)-\operatorname{ad}_{K m} K u+K \operatorname{ad}_{K u}^{*} m=0
\end{array}
$$

Thus the forward evolution of the cubic system is determined by the following set of four PDEs:

$$
\begin{aligned}
\dot{h}+(D h) K m & =0 \\
\dot{m}-u+\mathrm{ad}_{K m}^{*} m & =0 \\
\dot{u}-(D h)^{\top} p-K^{-1} \mathrm{ad}_{K m} K u+\mathrm{ad}_{K u}^{*} m & =0 \\
\dot{p}+\nabla \cdot(p \otimes K m) & =0
\end{aligned}
$$

Next we discuss how to use these shooting equations for the cubic evolution to define the cubic regression problem.

## Appendix C. Euler-Lagrange for shooting problem

Let, $J_{i}$, for $i=1 \ldots N$, denote $N$ measured images at timepoints, $t_{i} \in(0,1)$. Let us assume there are no measurements at the end points, i.e., neither at $t=0$, nor at $t=1$. Let $t_{c} \in(0,1)$, for $c=1 \ldots C$, denote $C$ data-independent fixed control locations. The control locations also implicitly define $C+1$ intervals or partitions in ( 0,1 ). Let us denote these intervals as $\mathcal{I}_{c}$, for $c=1 \ldots(C+1)$.

For regression on such a data configuration, the least-squares energy takes the form,
$E(h)=\frac{1}{2 \sigma_{I}^{2}} \sum_{c=1}^{C+1} \sum_{i \in \mathcal{I}_{c}} \mathrm{~d}^{2}\left(I^{0} \circ h\left(t_{i}\right), J_{i}\right)$,
subject to the dynamic constraints.
We first write the constrained energy minimization problem as,

$$
\begin{equation*}
\underset{h, m, u}{\operatorname{minimize}} \quad E(h) \tag{C.1}
\end{equation*}
$$

$$
\left.\begin{array}{rl}
\dot{h}+(D h) K m & =0 \\
\dot{m}-u+\mathrm{ad}_{K m}^{*} m & =0 \\
\dot{u}-(D h)^{\top} p-K^{-1} \mathrm{ad}_{K m} K u+\mathrm{ad}_{K u}^{*} m & =0 \\
\dot{p}+\nabla \cdot(p \otimes K m) & =0
\end{array}\right\} \text { for each }
$$

subject to continuity of $h, m$, and $u$ at $C$ joins.
The unconstrained Lagrangian for spline regression for a fixed initial image $I^{0}$ takes the form:

$$
\begin{aligned}
& \tilde{E}(h,\left.m, u, p, p^{t_{c}}, \lambda_{h}, \lambda_{m}, \lambda_{u}, \lambda_{p}\right) \\
&= \frac{1}{2 \sigma_{I}^{2}} \sum_{c=1}^{c+1} \sum_{i \in \mathcal{I}_{c}} \mathrm{~d}^{2}\left(I^{0} \circ h\left(t_{i}\right), J_{i}\right)+\frac{1}{2 \sigma_{u}^{2}} \int_{0}^{1}\langle u, K u\rangle_{L^{2}} d t \\
&+\sum_{c=1}^{c+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{h c}, \dot{h}_{c}+\left(D h_{c}\right) K m_{c}\right\rangle_{L^{2}} d t \\
&+\sum_{c=1}^{C+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{m c}, \dot{m}_{c}-u_{c}+\mathrm{ad}_{K m_{c}}^{*} m_{c}\right\rangle_{L^{2}} d t \\
& \quad+\sum_{c=1}^{c+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{u c}, \dot{u}_{c}-\left(D h_{c}\right)^{\top} p_{c}-K^{-1} \mathrm{ad}_{K m_{c}} K u_{c}+\mathrm{ad}_{K u_{c}}^{*} m_{c}\right\rangle_{L^{2}} d t \\
& \quad+\sum_{c=1}^{C+1} \int_{\mathcal{I}_{c}}\left\langle\lambda_{p c}, \dot{p}_{c}+\nabla \cdot\left(p_{c} \otimes K m_{c}\right)\right\rangle_{L^{2}} d t, \text { and }
\end{aligned}
$$

subject to continuity of $h, m$, and $u$ at $C$ joins.
The second term is the relaxation energy term, $\|u\|_{\mathfrak{q}^{*}}^{2}$. This term acts as a regularizer on the force along the full path in diffeomorphisms.

## C1. Variations

We discuss each piece of this optimization separately and combine the result in the end of this section. It is convenient to first derive it for the case with no controls, i.e., $C=0$. The unconstrained Lagrangian takes the form,
$\tilde{E}\left(h, m, u, p, \lambda_{h}, \lambda_{m}, \lambda_{u}, \lambda_{p}\right)=\frac{1}{2 \sigma_{I}^{2}} \sum_{i=1}^{N} \mathrm{~d}^{2}\left(I^{0} \circ h\left(t_{i}\right), J_{i}\right)$

$$
\begin{aligned}
& +\frac{1}{2 \sigma_{u}^{2}} \int_{0}^{1}\langle u, K u\rangle_{L^{2}} d t+\int_{0}^{1}\left\langle\lambda_{h}, \dot{h}+(D h) K m\right\rangle_{L^{2}} d t \\
& +\int_{0}^{1}\left\langle\lambda_{m}, \dot{m}-u+\mathrm{ad}_{K m}^{*} m\right\rangle_{L^{2}} d t \\
& +\int_{0}^{1}\left\langle\lambda_{u}, \dot{u}-(D h)^{\top} p-K^{-1} \mathrm{ad}_{K m} K u+\mathrm{ad}_{K u}^{*} m\right\rangle_{L^{2}} d t \\
& +\int_{0}^{1}\left\langle\lambda_{p}, \dot{p}+\nabla \cdot(p \otimes K m)\right\rangle_{L^{2}} d t
\end{aligned}
$$

The variations with respect to dual adjoint variables give the dynamic constraints back. We write the variations with respect to all the primals as:

$$
\begin{aligned}
\delta_{h} \tilde{E}= & \sum_{i=1}^{N}\left\langle\frac{1}{\sigma_{I}^{2}} \nabla_{h\left(t_{i}\right)} \mathrm{d}\left(I^{0} \circ h\left(t_{i}\right), J_{i}\right), \delta_{h}\left(t_{i}\right)\right\rangle \\
& +\left\langle\lambda_{h}(1), \delta h(1)\right\rangle-\left\langle\lambda_{h}(0), \delta h(0)\right\rangle \\
& +\int_{0}^{1}\left\langle-\dot{\lambda}_{h}+\nabla \cdot\left(p \otimes \lambda_{u}-\lambda_{h} \otimes K m\right), \delta h\right\rangle d t \\
\delta_{m} \tilde{E}= & \left\langle\lambda_{m}(1), \delta m(1)\right\rangle-\left\langle\lambda_{m}(0), \delta m(0)\right\rangle \\
& +\int_{0}^{1}\left\langle-\dot{\lambda}_{m}+K\left((D h)^{\top} \lambda_{h}\right)+\operatorname{ad}_{K m} \lambda_{m}-K \operatorname{ad}_{\lambda_{m}}^{*} m\right. \\
& \left.+K \operatorname{ad}_{K u}^{*} K^{-1} \lambda_{u}+\operatorname{ad}_{K u} \lambda_{u}-K\left(\left(D \lambda_{p}\right)^{\top} p\right), \delta m\right\rangle d t
\end{aligned}
$$

$$
\begin{aligned}
\delta_{u} \tilde{E}= & \left\langle\lambda_{u}(1), \delta u(1)\right\rangle-\left\langle\lambda_{u}(0), \delta u(0)\right\rangle \\
& +\int_{0}^{1}\left\langle-\dot{\lambda}_{u}+\frac{1}{\sigma_{u}^{2}} K u-\lambda_{m}-K \operatorname{ad}_{K m}^{*} K^{-1} \lambda_{u}-K \operatorname{ad}_{\lambda_{u}}^{*} m, \delta u\right\rangle d t \\
\delta_{p} \tilde{E}= & \left\langle\lambda_{p}(1), \delta p(1)\right\rangle-\left\langle\lambda_{p}(0), \delta p(0)\right\rangle \\
& +\int_{0}^{1}\left\langle-\dot{\lambda}_{p}-(D h) \lambda_{u}-\left(D \lambda_{p}\right) K m, \delta p\right\rangle d t
\end{aligned}
$$

At the optimum, the above must vanish. Thus, this results in the adjoint system presented in (11) for the general case with C controls.

The boundary conditions are,

$$
\begin{array}{ll}
\delta_{h(0)} E=-\lambda_{h}(0) & \delta_{h(1)} E=\lambda_{h}(1) \\
\delta_{m(0)} E=-\lambda_{m}(0) & \delta_{m(1)} E=\lambda_{m}(1) \\
\delta_{u(0)} E=-\lambda_{u}(0) & \delta_{u(1)} E=\lambda_{u}(1) \\
\delta_{p(0)} E=-\lambda_{p}(0) & \delta_{p(1)} E=\lambda_{p}(1)
\end{array}
$$

Notice that we can also rewrite the adjoint system in (11) for $C=0$ by removing the $K$ operator and using their duals, $v=K m$ and $f=K u$, and the conjugate operator, $\mathrm{ad}_{X}^{\dagger}$ is $\mathrm{ad}_{X}^{\dagger}=K \mathrm{ad}_{X}^{*} L$, such that:

$$
\begin{aligned}
& \dot{\lambda}_{h}-\nabla \cdot\left(p \otimes \lambda_{u}-\lambda_{h} \otimes v\right)=0 \\
& \dot{\lambda}_{m}-\operatorname{ad}_{v} \lambda_{m}+\operatorname{ad}_{\lambda_{m}}^{\dagger} v-\operatorname{ad}_{f}^{\dagger} \lambda_{u} \\
& \quad-\operatorname{ad}_{f} \lambda_{u}+K\left(\left(D \lambda_{p}\right)^{\top} p\right)-K\left((D h)^{\top} \lambda_{h}\right)=0 \\
& \dot{\lambda}_{u}-\frac{1}{\sigma_{u}^{2}} f+\lambda_{m}+\operatorname{ad}_{v}^{\dagger} \lambda_{u}+\operatorname{ad}_{\lambda_{u}}^{\dagger} v=0 \\
& \dot{\lambda}_{p}+(D h) \lambda_{u}+\left(D \lambda_{p}\right) v=0
\end{aligned}
$$

## C1.1. For the data fit constraints

The gradient of the data match term with respect to $h$ is:

$$
\begin{aligned}
& \frac{1}{2 \sigma_{I}^{2}}\left\langle\nabla_{h\left(t_{i}\right)} \mathrm{d}\left(I^{0} \circ h\left(t_{i}\right), J_{i}\right), \delta_{h}\left(t_{i}\right)\right\rangle \\
& \quad=\frac{1}{\sigma_{I}^{2}}\left\langle\left(I^{0} \circ h\left(t_{i}\right)-J_{i}\right), \nabla_{h\left(t_{i}\right)}\left(I^{0} \circ h\left(t_{i}\right)\right) \delta_{h}\left(t_{i}\right)\right\rangle \\
& \quad=\frac{1}{\sigma_{I}^{2}}\left\langle\left(I^{0} \circ h\left(t_{i}\right)-J_{i}\right), \nabla I^{t_{i}} \delta_{h}\left(t_{i}\right)\right\rangle
\end{aligned}
$$

Note there is no application of the chain rule in the last step above for the term, $\nabla_{h\left(t_{i}\right)}\left(I^{0} \circ h\left(t_{i}\right)\right)$ because the gradient is computed with respect to $h\left(t_{i}\right)$. This is equivalent to the substitution of $h\left(t_{i}\right)$ as $y$, which then is equivalent to taking the derivative of $I(y)$ with respect to $y$. Thus, this gradient of the data fit constraints with respect to $h$, results in jumps during the backward integration of the adjoint variable, $\lambda_{h}$, as,
$\lambda_{h}\left(t_{i}^{+}\right)-\lambda_{h}\left(t_{i}^{-}\right)=\frac{1}{\sigma_{I}^{2}}\left(I^{0} \circ h\left(t_{i}\right)-J_{i}\right) \nabla I^{t_{i}}$.

## C1.2. For the continuity constraints

We derive the variations of the energy functional for the joins and study the continuity of the adjoint system in the interval $(0,1)$. For this analysis, we introduce again the subscripts $c$ to denote the intervals. We first rewrite the above functional here as,
$\tilde{E}=$ fit of the data within each interval, $\mathcal{I}_{c}$

+ dynamics within each interval, $\mathcal{I}_{c}$
$+\sum_{c=1}^{C}\left\langle\alpha_{h c}, h_{c}\left(t_{c}\right)-h_{c+1}\left(t_{c}\right)\right\rangle_{L^{2}}$
$+\sum_{c=1}^{c}\left\langle\alpha_{m c}, m_{c}\left(t_{c}\right)-m_{c+1}\left(t_{c}\right)\right\rangle_{L^{2}}$
$+\sum_{c=1}^{c}\left\langle\alpha_{u c}, u_{c}\left(t_{c}\right)-u_{c+1}\left(t_{c}\right)\right\rangle_{L^{2}}$

Notice that the Lagrangian adjoint variables, $\alpha_{h c}, \alpha_{m c}$ and $\alpha_{u c}$, do not vary with time. We write the variations with respect to all the primals at joins as:

$$
\begin{aligned}
\delta_{h c\left(t_{c}\right)} \tilde{E} & =\lambda_{h c}\left(t_{c}\right)+\alpha_{h c} \\
\delta_{h c+1\left(t_{c}\right)} \tilde{E} & =-\lambda_{h c+1}\left(t_{c}\right)-\alpha_{h c} \\
\delta_{m c\left(t_{c}\right)} \tilde{E} & =\lambda_{m c}\left(t_{c}\right)+\alpha_{m c} \\
\delta_{m c+1\left(t_{c}\right)} \tilde{E} & =-\lambda_{m c+1}\left(t_{c}\right)-\alpha_{m c} \\
\delta_{u c\left(t_{c}\right)} \tilde{E} & =\lambda_{u c}\left(t_{c}\right)+\alpha_{u c} \\
\delta_{u c+1\left(t_{c}\right)} \tilde{E} & =-\lambda_{u c}\left(t_{c}\right)-\alpha_{u c}
\end{aligned}
$$

Equating all the above variations to zero and algebraically eliminating variables, we get,
$\lambda_{h c}\left(t_{c}\right)=\lambda_{h c+1}\left(t_{c}\right)$
$\lambda_{m c}\left(t_{c}\right)=\lambda_{m c+1}\left(t_{c}\right)$
$\lambda_{u c}\left(t_{c}\right)=\lambda_{u c+1}\left(t_{c}\right)$
This proves that $\lambda_{h c}, \lambda_{m c}$ and $\lambda_{u c}$ are continuous at the boundaries of the control point locations.

## Supplementary materials

Supplementary material associated with this article can be found, in the online version, at 10.1016/j.media.2015.04.012.

## References

Adams, J.F., 1969. Lectures on Lie Groups. University of Chicago Press.
Amit, Y., Grenander, U., Piccioni, M., 1991. Structural image restoration through deformable templates. J. Am. Stat. Assoc. 86 (414), 376-387.
Arnol'd, V.I., 1966. Sur la géométrie différentielle des groupes de Lie de dimension infinie et ses applications à l'hydrodynamique. Ann. Inst. Fourier 16, 319-361.
Beg, M.F., Miller, M.I., Trouvé, A., Younes, L., 2005. Computing large deformation metric mappings via geodesic flows of diffeomorphisms. Int. J. Comput. Vis. 61 (2), 139157.

Bruveris, M., Gay-Balmaz, F., Holm, D., Ratiu, T., 2011. The momentum map representation of images. J. Nonlinear Sci. 21 (1), 115-150. doi:10.1007/s00332-010-9079-5.
Byrd, R., Lu, P., Nocedal, J., Zhu, C., 1995. A limited memory algorithm for bound constrained optimization. SIAM J. Sci. Comput. 16 (5), 1190-1208. doi:10.1137/0916069.
Camarinha, M., Leite, F.S., Crouch, P., 1995. Splines of class ck on non-euclidean spaces. IMA J. Math. Control Inform. 12 (4), 399-410.
Chambolle, A., 2004. An algorithm for total variation minimization and applications. J. Math. Imaging Vis. 20 (1-2), 89-97. doi:10.1023/B:JMIV.0000011325.36760.1e.
Chevalley, C., 1999. Theory of Lie Groups: 1, vol. 1. Princeton University Press.
Crouch, P., Leite, F.S., 1995. The dynamic interpolation problem: on riemannian manifolds, lie groups, and symmetric spaces. J. Dyn. Control Syst. 1 (2), 177-202.
Curtis, C.W., Reiner, I., 1962. Representation Theory of Finite Groups and Associative Algebras, vol. 356. AMS Bookstore.
Davis, B.C., 2008. Medical Image Analysis via Fréchet Means of Diffeomorphisms. ProQuest.
Davis, B.C., Fletcher, P.T., Bullitt, E., Joshi, S., 2010. Population shape regression from random design data. Int. J. Comput. Vis. 90 (2), 255-266.
Durrleman, S., Pennec, X., Trouv, A., Braga, J., Gerig, G., Ayache, N., 2013. Toward a comprehensive framework for the spatiotemporal statistical analysis of longitudinal shape data. Int. J. Comput. Vis. 103 (1), 22-59. http://dx.doi.org/10.1007/s11263-012-0592-x.

Fulton, M.W., Harris, J., 1991. Representation Theory: A First Course, 129. Springer.
Gay-Balmaz, F., Holm, D.D., Meier, D.M., Ratiu, T.S., Vialard, F.-X., 2012. Invariant higherorder variational problems II. J. Nonlinear Sci. 22 (4), 553-597.
Grenander, U., Miller, M.I., 1998. Computational anatomy: an emerging discipline. Q. Appl. Math. LVI (4), 617-694.
Gu, X., He, Y., Qin, H., 2006. Manifold splines. Graphical Models 68 (3), 237-254 http://www.sciencedirect.com/science/article/pii/S152407030600021X, http://dx.doi.org/10.1016/j.gmod.2006.03.004.
Hinkle, J., Fletcher, P., Joshi, S., 2014. Intrinsic polynomials for regression on Riemannian manifolds. J. Math. Imaging Vis. 50 (1-2), 32-52. doi:10.1007/s10851-013-0489-5.
Krakowski, K.A., 2003. Geometrical Methods of Inference. University of Western Australia.
Lorenzi, M., Ayache, N., Frisoni, G., Pennec, X., et al., 2010. 4d registration of serial brain's MR images: a robust measure of changes applied to alzheimer's disease. In: Spatio Temporal Image Analysis Workshop (STIA), MICCAI. Citeseer.
Machado, L., Leite, F.S., Hüper, K., 2006. Riemannian means as solutions of variational problems. LMS J. Comput. Math 9, 86-103.
Machado, L., Silva Leite, F., Krakowski, K., 2010. Higher-order smoothing splines versus least squares problems on riemannian manifolds. J. Dyn. Control Syst. 16 (1), 121148. http://dx.doi.org/10.1007/s10883-010-9080-1.

Miller, M.I., 2004. Computational anatomy: shape, growth, and atrophy comparison via diffeomorphisms. NeuroImage 23, 19-33.
Miller, M., Banerjee, A., Christensen, G., Joshi, S., Khaneja, N., Grenander, U., Matejic, L., 1997. Statistical methods in computational anatomy. Stat. Methods Med. Res. 6 (3), 267-299.
Niethammer, M., Huang, Y., Vialard, F.-X., 2011. Geodesic regression for image timeseries. In: Fichtinger, G., Martel, A., Peters, T. (Eds.), MICCAI 2011, LNCS, vol. 6892. Springer, Berlin/Heidelberg, pp. 655-662. doi:10.1007/978-3-642-23629-7_80.
Noakes, L., Heinzinger, G., Paden, B., 1989. Cubic splines on curved spaces. IMA J. Math. Control Inform. 6 (4), 465-473.
Radau, P., Lu, Y., Connelly, K., Paul, G., Dick, A., Wright, G., 2009. Evaluation framework for algorithms segmenting short axis cardiac MRI. MIDAS. http://hdl.handle.net/10380/3070.
Rudin, W., 1986. Real and Complex Analysis, third ed. New York: McGraw-Hill Inc.
Rudin, W., 1991. Functional Analysis. International Series in Pure and Applied Mathematics. McGraw-Hill, Inc., New York.
Schwartz, E., Jakab, A., Kasprian, G., Zllei, L., Langs, G., 2015. A locally linear method for enforcing temporal smoothness in serial image registration. In: Durrleman, S., Fletcher, T., Gerig, G., Niethammer, M., Pennec, X. (Eds.), Spatio-temporal Image Analysis for Longitudinal and Time-Series Image Data. In: Lecture Notes in Computer Science. Springer International Publishing, pp. 13-24. http://dx.doi.org/ 10.1007/978-3-319-14905-9_2.

Singh, N., Fletcher, P., Preston, J.S., King, R.D., Marron, J.S., Weiner, M.W., Joshi, S., 2014. Quantifying anatomical shape variations in neurological disorders. Med. Image Anal. 18 (3), 616-633. doi:10.1016/j.media.2014.01.001.
Singh, N., Hinkle, J., Joshi, S., Fletcher, P., 2013a. A hierarchical geodesic model for diffeomorphic longitudinal shape analysis. In: IPMI, vol. 7917, pp. 560-571.
Singh, N., Hinkle, J., Joshi, S., Fletcher, P., 2013b. A vector momenta formulation of diffeomorphisms for improved geodesic regression and atlas construction. In: ISBI, pp. 1219-1222. doi:10.1109/ISBI.2013.6556700.
Singh, N., Niethammer, M., 2014. Splines for diffeomorphic image regression. In: Golland, P., Hata, N., Barillot, C., Hornegger, J., Howe, R. (Eds.), Medical Image Computing and Computer-Assisted Intervention (MICCAI 2014). Lecture Notes in Computer Science, vol. 8674. Springer International Publishing, pp. 121-129. http://dx.doi.org/10.1007/978-3-319-10470-6_16.
Thompson, D.W., et al., 1942. On Growth and Form. Cambridge Univ. Press.
Thompson, P.M., Toga, A.W., 2002. Computing and Visualization in Science. A Framework for Computational Anatomy 5 (1), 13-34. doi:10.1007/s00791-002-0084-6.
Trouvé, A., Vialard, F.-X., 2012. Shape splines and stochastic shape evolutions: a second order point of view. Q. Appl. Math. 70 (2), 219-251.
Vercauteren, T., Pennec, X., Perchant, A., Ayache, N., 2009. Diffeomorphic demons: efficient non-parametric image registration. NeuroImage 45 (1), S61-S72.
Vialard, F.-X., Risser, L., Rueckert, D., Cotter, C., 2012. Diffeomorphic 3D image registration via geodesic shooting using an efficient adjoint calculation. Int. J. Comput. Vis. 97 (2), 229-241. doi:10.1007/s11263-011-0481-8.
Younes, L., 2010. Shapes and Diffeomorphisms, vol. 171. Springer, Berlin.
Younes, L., Arrate, F., Miller, M.I., 2009. Evolution equations in computational anatomy. NeuroImage 45 ( 1 Suppl.), S40-S50.


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