Abstract

We introduce a volumetric space-time technique for the reconstruction of moving and deforming objects from point data. The output of our method is a four-dimensional space-time solid, made up of spatial slices, each of which is a three-dimensional solid bounded by a watertight manifold. The motion of the object is described as an incompressible flow of material through time. We optimize the flow so that the distance material moves from one time frame to the next is bounded, the density of material remains constant, and the object remains compact. This formulation overcomes deficiencies in the acquired data, such as persistent occlusions, errors, and missing frames. We demonstrate the performance of our flow-based technique by reconstructing coherent sequences of watertight models from incomplete scanner data.

CR Categories: I.3.7 [Computer Graphics]: Three-Dimensional Graphics and Realism

Keywords: Reconstruction, space-time, volumetric techniques

1 Introduction

Recent advanced scanning technologies together with increasing computational power allow the space-time capture of three-dimensional objects as they move and deform. Systems such as [Zhang et al. 2003; Fong and Buron 2005; Koninckx and van Gool 2006] produce dense point samples over large parts of the surface of a moving object, at rates from ten to thirty frames per second. As these technologies mature, they make it possible to capture and reconstruct both the deforming model and its motion. Organizing the data captured by these scanners into a coherent model of a deforming object is a challenging computational problem that is just beginning to be addressed [Mitra et al. 2007; Wand et al. 2007; Pekelny and Gotsman 2008].

As pointed out in one of the first applications of real-time scanning [Rusinkiewicz et al. 2002], the main challenge posed by the time component is to fill in missing data by accumulating information over time. The input scans are typically collected by a small set of synchronized cameras. Because of the small number of fixed views, large parts of the surface are occluded in each frame, leading to gaping holes that often persist across many frames. Current real-time systems also suffer from low resolution, insufficient frame rate and noise. While it is reasonable to assume advances in real-time scanning technology will improve the data quality, persistent occlusions will always be an issue.

To complete the missing data in a principled way we take advantage of space-time coherence and adopt a global approach, which considers all frames simultaneously. Furthermore, to guide the reconstruction, we include as many reasonable physical assumptions as we can into our computation. First, we directly reconstruct a deforming solid, that is, a four-dimensional space-time surface. By explicitly modeling the mass field of the object, we leverage the knowledge that the boundary of the reconstructed object is a watertight manifold. Most importantly, we explicitly model the flow of this material through time. We constrain all of the material in
one time-frame to move to some nearby position in the next, ensuring mass conservation, while preventing mass from concentrating or dispersing, producing what we call an incompressible flow. We also introduce a momentum term which ensures that material moves smoothly through time. We use this flow terminology somewhat loosely; as will be seen, our method does not produce a physically accurate fluid flow; nor does it need to.

This incompressible flow prior puts a very strong constraint on the shape of the four-dimensional solid and on how its 3D spatial slices evolve over time. Using all the frames simultaneously to compute the flow, data from arbitrarily distant frames is used to plausibly complete missing data in occluded regions. We can also successfully reconstruct entirely missing frames by extrapolating information from sampled ones, overcoming sparse frame rates (Figure 6).

Our formulation requires the fairly mild assumptions that the object moves smoothly through time and that the speed of movement between consecutive time frames is bounded. We do not need to assume that the motion is globally or locally rigid, nor do we have any assumptions on the topology or geometry of the object. In contrast to many previous methods [Marschner et al. 2000; Guskov et al. 2003; White et al. 2007] we do not require additional space-time coherence information such as marker correspondences.

Specifically, each three-dimensional frame is represented by its characteristic function on a regular grid. We construct the mass field, i.e. the characteristic function, and its flow simultaneously for all frames. To do so, we constrain all the material at time \( t \) to move to some adjacent grid cell at time \( t + 1 \). We set the material in any cell identified as inside the object to be one and in any outside cell to be zero, avoiding compression or dilation inside each cell. The estimation of both distribution of material in space and its flow is computed by iteratively updating and solving a linear system, applying at each iteration a simple re-weighting step to enforce incompressibility and enhance the object boundaries.

The obvious difficulty with our approach is that the entire volume and its flow are explicitly represented, producing a very large linear system. We handle this by employing a preconditioned Krylov subspace iterative solver that exploits the sparsity and the specific spectral properties of the system.

2 Background

While numerous methods exist for reconstructing static 3D models, none of them is directly applicable to the dynamic setting. The main challenge faced by dynamic reconstruction is to properly define the relationship between the temporal and spatial dimensions in order to accumulate information over time to correctly complete locally missing data [Grosso et al. 1989; Rusinkiewicz et al. 2002]. Much of the attention in 4D reconstruction has been paid to processing specific shapes such as humans or garments where additional information is available to guide the reconstruction, e.g., marker correspondences between scans [Marschner et al. 2000; Guskov et al. 2003; White et al. 2007] or between the scans and a template [Allen et al. 2002; Anguelov et al. 2004; Zhang et al. 2004; Anguelov et al. 2005]. Unmarked data is much simpler to acquire making a markerless approach like ours more general.

Arbitrary shape reconstruction from point cloud sequences is usually treated as a surface deformation over time. Specifically, Shinya [2004] extracts an initial surface from the first point cloud, and evolves it towards the subsequent point clouds while minimizing a deformation energy computed over the surface triangulation. Wang et al. [2005] further refine this energy using harmonic maps. Anuar and Guskov [2004] use optical flow to estimate the deformation, switching from surface representation to volumetric flow integration. Recently, Pekelny and Gotsman [2008] introduce another method to accumulate information for the specific case of articulations of rigid parts. The input point cloud is manually segmented and the parts are tracked over time assuming that each frame is a good initial guess for an iterative closest point rigid motion registration with the next frame. These approaches are sensitive to the quality of the initial surface since they accumulate information only forward in time.

To integrate time information for arbitrary shapes, Wand et al. [2007] and Mitra et al. [2007] formulate the reconstruction problem directly in 4D, focusing on inter-frame registration. Wand et al. [2007] optimize a 4D shape represented by a set of surfels. They define the surface from statistical densities, imposing spatial smoothness and rigid motion priors. Mitra et al. [2007] register point clouds directly on a 4D hypersurface, assuming rigid inter-frame motion. With dense temporal sampling, this allows estimating local deformations of the model. We approach the problem in 4D as well, but by operating on a volume, rather than a surface like Wand et al. [2007] or Mitra et al. [2007]. In particular, we generate a sequence of watertight manifolds as output, and we model the flow of mass in order to compensate for missing data. Unlike these techniques our method does not require dense spatial sampling to complete missing data.

Other approaches address the related problem of reconstructing moving objects from video sequences [Magnor and Goldlücke 2004; Goldlücke and Magnor 2005; Goldlücke et al. 2007]. They carve a 4D hypersurface and then optimize it by enforcing photo-consistency with a 2D video sequence.

The ability to reconstruct a watertight object using the strong incompressible flow prior distinguishes our method from previous techniques. This prior leads to more reliable reconstruction enabling us to obtain accurate, watertight reconstructions from poorly sampled data.

As a last note, our solution to the 4D problem uses the concept of material flow. Similar techniques have been devised for 2D video in the Computer Vision literature [Ullman 1979; Gupta and Kanal 1995; Vedula et al. 1999; Barron and Thacker 2004], to track motions through minimization of specific image similarity measures.

3 Problem Formulation

We consider the problem of reconstructing a moving and possibly deforming object of arbitrary topology given a sequence of three-dimensional frames. Each frame consists of a cloud of points in 3D sampled over the object’s surface. Our goal is to reconstruct a watertight surface in space-time, that is, a three-dimensional surface...
embedded in 4D. We represent this 4D solid using its characteristic function on a 4D grid, from which we extract the actual object surface at a final step (Section 4.3).

For the sake of simplicity we describe our formulation for an object in one-dimensional space deforming through time. The domain is then an \((n + 1) \times (m + 1)\) space-time grid, and we use \(i, j\) for space indices and \(t\) for time ones. The three- and four-dimensional formulations we use later are straightforward generalizations obtained by replacing the spatial adjacency relationships used in one dimension with those for higher dimensions (Figure 2). The characteristic function values \(x_t^j\) at each cell can be seen as representing the amount of material in the cell. Using this representation we describe the motion and deformation of the object over time as flow of material through space-time. The flow is represented by variables \(v_{i,j}^t\) representing the amount of material moving from cell \(x_t^j\) at time \(t\) to another cell \(x_{t+1}^j\) at time \(t+1\). Note that, in contrast to the simulation of fluid flows, material in our model moves through time at a constant rate, so that higher flow values on an edge mean more material moving through the edge. We formulate the reconstruction as a solution to a constrained minimization problem, based on the following set of assumptions.

**Flow incompressibility:** We interpret flow incompressibility as constraining the amount of material in any cell to be equal to the amount of material flowing into the cell from the previous time frame, and also to the amount of material flowing out of the cell into the next time frame. Using the previous notation, this is:

\[
x_t^j = \sum_j v_{i,j}^{t-1}, \quad i = 0 \ldots n, \quad t = 1 \ldots m \tag{1}
\]

\[
x_t^j = \sum_j v_{i,j}^t, \quad i = 0 \ldots n, \quad t = 0 \ldots m - 1 \tag{2}
\]

**Bounded speed:** We expect the grid resolution in time to be sufficiently dense with respect to the deformation speed. Specifically, we assume that at each time step material can only move to temporally adjacent cells (shown in Figure 2). In \(1D \times time\) this means that \(v_{i,j}^t\) is non-zero only if \(i-1 \leq j \leq i+1\). This constraint is handled implicitly by including in the formulation only the variables for \(v_{i,j}^t\), that can potentially be non-zero. Note that the actual sampling of the data in time could be at a lower resolution than that of the solution grid, or in other words we can have time-frames with no samples.

**Spatial continuity:** In the characteristic function, we expect values \(x_t^j\) in spatially adjacent cells to be identical everywhere, except across the object boundaries. Because the scans give an incomplete representation of the surface, however, we do not know where some of the boundaries are and in addition we cannot be sure of the in/out orientation at all known boundaries. We therefore require the values of \(x_t^j\) to be constant everywhere except at a small number of sharp discontinuities. Minimizing a functional such as

\[
F_c(x_t^j) = \sum_{i=1}^m \left(x_t^j - x_{t-1}^j\right)^{0.8}
\]

(3)

gives this effect. The sub-linear exponent 0.8 in the norm penalizes many small discontinuities more than a few large ones [Levin et al. 2007]. It would be easier, computationally, to use a least-squares error functional, since the derivative would be linear. But that would have exactly the wrong effect; with an exponent of 2 instead of 0.8, the minimum would be achieved when the differences between cell values \(x_t^j\) are evenly distributed across all cells. We address this computational issue, and also some refinements of this functional, in Section 4.2.

Note that \(F_c\) only links cells adjacent to one another in the spatial dimension. We use the flow to control the variation of the function in the time dimension.

**Flow momentum:** We expect the object deformation, and hence the flow, to be smooth in time. This smoothness is captured by a momentum term that penalizes the flow for changing direction from one time frame to the next:

\[
F_m(v_{i,j}) = \sum_{i,j=0}^{m-1} (v_{i,j} - v_{i+1,j})^2
\]

(4)

This term uses the discretized flow directions available on the grid, which is not correct at the object boundaries; there we rely on the spatial continuity term to enforce continuity of momentum as well. We tried including a similar boundary-enforcement mechanism in the momentum term, but we found that the slight improvement in performance did not justify the additional cost.

Figure 3: Reconstruction of a scanned moving hand puppet. Left: Input scan points to the flow solver (green points), with the initial inside cell labeling (dark blue cells). Center: Two flow solver iterations: mass cells is represented by a grayscale map. Cells determined to be inside by the solver are colored dark blue. Right: Our final reconstruction.
Data fidelity: The values of the characteristic function $x$ for some of the grid cells are known even before the optimization starts. Specifically, we assign $x_i = 1$ to cells containing scan points, and we mark those as boundary cells. Values of zero and one are assigned to additional cells, both at initialization and during the computation, as discussed in Section 4. These values are treated as hard constraints and handled implicitly by using back-substitution, moving the corresponding variables from the system.

Problem formulation: The solution to our space-time reconstruction problem is computed by optimizing a weighted combination of spatial continuity $F_c$ and flow momentum $F_m$, $F = \alpha F_c + (1 - \alpha) F_m$, subject to the incompressibility constraints. We used $\alpha = 2/3$ for the 2D×time examples, and $\alpha = 1/3$ for the 3D×time ones, giving higher weight to flow smoothness. This difference of weighting was necessary to compensate for the increased number of spatially adjacent cells in each time frame. We observed that when there was a lot of missing data, putting more weight on flow momentum filled in missing areas faster, accelerating convergence.

4 Solution Mechanism
4.1 Preprocessing and Initialization
The characteristic function values for some cells can be reliably computed without the flow computation. Initializing as many cells as possible drastically reduces the number of variables in the functional to be optimized, significantly increasing solution speed. It also improves the solution accuracy. Therefore we do as much preprocessing as we can before attempting the flow computation.

We begin by computing a low-resolution visual hull for each frame, based on the scan planes and the scanner positions and orientations. This allows us to loosely label many outside cells. Certainly in the presence of noise and holes, estimation of the visual hull is error-prone, nevertheless, at this preprocessing step high precision is not required.

We select a small number of cells to loosely label as inside using the following heuristic. We compute for each frame a low-resolution unsigned distance field measuring the distance from scan data, and select cells that are both local maxima of the distance function and are inside the visual hull. The idea is that such cells simultaneously lie near the medial axis of the surface and inside the visual hull, and as such are very likely to be inside the object itself.

This is followed by a low-resolution 4D surface reconstruction, using the FEM reconstruction method of Sharf et al. [2007]. This algorithm takes as input both the data points and the above described inside/outside labels. It computes a smooth function that is negative inside the object and positive outside. At such low resolution, this computation is very efficient. We utilize the FEM solution in two ways. First, we use it to improve our inside/outside labeling; we label maxima of the FEM function as inside the object and its minima as inside. Second, we use it to assign normals to the data points. We use these normals to modify the spatial continuity functional $F_c$, as described below in Section 4.2. Examples of initial cell labeling are shown in Figure 3.

4.2 Optimization
We now describe our optimization of the flow functional. We use an Iteratively Reweighted Least Squares [Holland and Welsch 1977] approach to handle the sub-linear exponent in $F_c$. We introduce a weight $w_{i,j}$ for every pair of spatially adjacent cells $x_i$ and $x_j$, which is used to modify the spatial continuity functional $F_c$. The overall computation requires iteratively solving a large linear system, modifying the $w_{i,j}$ at each iteration.

\[ F_c(x_i) = \sum_{i=1}^{n} \sum_{j=1}^{n} w_{i,j} (x_i - x_j)^2 \] (5)

where the initial value of $w_{i,j}$ is zero if the face $(i,j)$ is an outward boundary face, and one everywhere else. Due to the subsequent global optimization, local normal estimation errors have no significant impact on the final result. In figure 4 we show that mass propagates from the small initial guess (left) to the actual mass (right) due to this continuity setting.

Iterations: After the first step the resulting distribution of $x'_i$ values provides some indication of where discontinuities are likely to arise; the greater the discontinuity between two cells in the current solution, the more likely a boundary falls there in the final result.

Iteratively Reweighted Least Squares [Holland and Welsch 1977] suggests a principled way to use this information: we set the weight $w_{i,j}$ in Equation (5) to be $w_{i,j} = (x_i' - x_j')^{0.8 - 2}$, using the values of $x$ from the previous iteration. This effectively guarantees that if the scheme converges, the solution we have is the minimum of the original functional $F_c$. To normalize the weights, we bound them from above by 0.0010.8 - 2 and scale into the range (0, 10). As expected, as the iteration proceeds the values of $x_i'$ become concentrated at zero and one as desired (Figure 6).

Regularization: The matrix in the linear system we solve at each iteration is of the form of a discrete Laplacian, and as such can be ill-conditioned, causing numerical instability. We use a standard regularization technique [Hansen 1998] adding the following term to our functional:

\[ F_r = \sum (x_i')^2 + \sum (x_i')^2. \]

$F_r$ is assigned a fairly small weight of 0.0025, since while we want the regularization to stabilize the system, it should have little effect on the final result.

Bootstrapping and clamping: As the flow solution is computed iteratively, many of the values obtained for $x_i'$ approach the extrema of zero and one; in fact, the linear system may also produce values of $x_i'$ which are negative or greater than one. In our experiments only slight deviations occurred from the [0–1] range. We use these extreme values of $x_i'$ to classify the corresponding cells as inside or outside. In subsequent iterations we clamp the values at these cells

Figure 4: Graphs demonstrating mass incompressibility and spatial continuity principles on the Elephant data set (video). Left: Our initial guess labels only few cells as inside mass per frame. Right: Spatial continuity term propagates the initial mass throughout the solid, while incompressibility term keeps total mass nearly level across frames.
Figure 5: A rotating 2D boomerang sweeps out the space-time surface (a); green points show where the surface was sampled. A horizontal slice represents the state of the boomerang at one time frame. Comparing 3D RBF reconstruction (b), direct FEM reconstruction (c), our flow reconstruction (d), we note how our flow overcomes the unsampled frames and preserves the boomerang’s concavity.

to be exactly one or zero, and use them as additional data fidelity constraints, using back-substitution.

Clamping introduces some rounding error to our mass conservation, but the error is essentially random and does not lead to a serious bias (around 2%, as seen in Figure 4). It also enforces incompressibility; an $x_i^t$ greater than one would represent a concentration of material in a cell, which is not allowed in our flow formulation.

Our formulation also allows us to enforce some flow constraints via back-substitution, further reducing the number of variables in the system. Specifically, given a cell with value zero, we can trivially constrain all flow incoming into the cell or outgoing from it to zero. Similarly, if the incoming or outgoing flow for a cell sums to zero, the cell value is set to zero.

The algorithm effectively converges when the values for all the cells are set by the bootstrapping. In practice, to speed up the process, we typically stop once 90% to 95% of the cell values are set, with the final step of the algorithm (Section 4.3) resolving the classification for those based on smoothness considerations.

4.3 Postprocessing

In the postprocessing step, we refine the optimal flow solution to generate the final high resolution output surfaces. This step serves two purposes; first, it improves the flow resolution if necessary, and second, it replaces the characteristic function with an estimated signed distance function, which is easier to contour. The refinement is done on a hexadeca-tree (4D octree) so that we can increase resolution near the surface to account for fine details in the input data.

We use a standard least squares solver for a Laplace operator on the hexadeca-tree, which maximizes local space-time smoothness. The in/out voxel labels of the flow solution are used as boundary conditions in the Laplace system. We assign signed distance values from the surface to cells in this tree based on both the flow solution and distance to the input data points. Cells deep within the object and cells containing data points are weighted more heavily than cells near the surface but with no data. Thus, the space close to the surface, is initialized as unknown.

The Laplace solution defines a 4D implicit function whose zero set represents the desired 3D surface across time. To extract polygonal surfaces, a standard marching cubes surface extraction is applied at each temporal cross-section of the implicit field independently.

5 Solving the Linear System

To utilize the presented formulation we must efficiently handle the large linear systems that arise during the computation. In this section we show how we exploit the matrix block structure to compute a solution quickly and with a minimal overhead in terms of memory. To accomplish this, we use an augmentation approach [Golub and Greif 2003].

At each iteration our formulation yields linear systems of the form

$$\begin{pmatrix} A & B^T \\ B & 0 \end{pmatrix} \begin{pmatrix} x \\ v \end{pmatrix} = \begin{pmatrix} f \\ g \end{pmatrix}$$

where $A$ arises from the optimized functional and $B$ reflects the constraints (Section 3). The matrix is large, sparse and indefinite, and standard direct techniques based on forming the $LDL^T$ decomposition with symmetric pivoting result in significant loss of sparsity. This in turn would lead to unreasonably large memory requirements and prohibitive computational cost. Therefore, we use the MINRES iterative method with a symmetric positive-definite block diagonal preconditioner which is well suited to the problem and requires a minimal overhead.
We set the preconditioner to be

\[ M = \begin{pmatrix} A + \gamma^{-1}B^T B & 0 \\ 0 & \gamma I \end{pmatrix}, \]

where \( \gamma \) is a scalar and \( I \) is the identity matrix. Denote the matrix of the system by \( K \). The matrix \( A + \gamma^{-1}B^T B \), which is a stabilized primal Schur complement of \( K \), is sparse and symmetric positive-definite. It can be shown that the preconditioned matrix \( M^{-1}K \) has an eigenvalue 1 of algebraic multiplicity equal to the dimension of \( A \). Furthermore, the negative eigenvalues of the preconditioned matrix are all strictly between \(-1\) and 0, and in our setting most of them are located near \(-1\), since nearly zero eigenvalues of \( A \) are mapped into eigenvalues of the preconditioned matrix that are nearly \(-1\) [Greif and Schötzau 2006].

This high algebraic multiplicity is crucial, since the speed of convergence of MINRES (and in fact any preconditioned Krylov subspace method) primarily depends on how well the eigenvalues of the preconditioned matrix are clustered; in this case, we will be able to obtain convergence within a number of iterations which is significantly smaller than the dimensions of the linear system, and in fact also smaller than the number of constraints.

Our numerical solution is based on inner/outer iterations. In solving for \( A + \gamma^{-1}B^T B \) (the inner iteration) we apply a conjugate gradient solver, preconditioned with an incomplete Cholesky factorization with a drop tolerance of 0.01. Since the discrete operators \( A \) and \( B^T B \) have spectral norms of approximately the same order, we set \( \gamma = 1 \) throughout the computation. We solve the inner iterations with a convergence tolerance of \( 10^{-6} \). The outer MINRES iterations are solved with a convergence tolerance of 0.0001.

The use of a high drop tolerance for the incomplete factorization and a loose outer convergence tolerance make the solver computationally inexpensive, with small iteration counts and modest memory requirements. Even in 4 dimensions the solver effectively converges within fewer than a hundred outer iterations. Our solver implementation combines the TAUCS library [Toledo 2003] with our own specialized MINRES solver. We can solve the system either on a standard workstation (2GB RAM 2.4 GHz CPU) or very quickly on a small 40-cores cluster, as described in Section 6. As a comparison, a direct parallel LDL^T solver for the same problem takes more than a day to compute the solution on the same cluster.

6 Results

We have tested our method on several real and synthetic examples, in both 2D×time and 3D×time. In all cases, our method produced watertight surfaces with good quality completion of missing data. We have tested our method on several real and synthetic examples, in both 2D×time and 3D×time. In all cases, our method produced watertight surfaces with good quality completion of missing data. 

2D×time results. To illustrate the merits of the flow solution, we use some 2D×time examples. In Figures 5 and 6, we show a two-dimensional “boomerang shaped” object moving through time. We leave large portions of the moving object un-sampled in many frames, mimicking the effects of occlusion. Furthermore, in order to explore our method’s limits and robustness, we leave some frames completely empty (Figure 6 bottom row). Observe at the center of Figure 5 that the reconstructions generated by usual surface reconstruction, such as RBF [Ohtake et al. 2003] or direct FEM [Sharf et al. 2007], introduce large cavities where significant data is missing. We note that while these might be valid reconstructions of a 3D solid, they are inconsistent under our mass incompressibility assumption. Figure 6 illustrates the flow solution across several frames for the boomerang example. The areas of the final 2D cross sections are roughly equal across all frames; they are not perfectly identical since the clamping of material values to either zero or one at most voxels leads to rounding errors.

A similar behavior can be observed in Figures 7 and 8, which show a two-dimensional rounded box that splits from one connected component into two. Although there are very large chunks of missing data, our method faithfully reconstructs the geometry. In Figure 6, we again compare the results of our method on three other moving data-sets, whose statistics are reported in Table 1. For the flow computation, the spatial resolution of all of our frames is 64^3. For the surface extraction, the final resolution of the hexadeca-tree varies from 64^3 to 256^3 depending on the model smoothness. While this resolution is not particularly fine, it was sufficient for the purpose of identifying inside/outside cells in our tests.

In our computations we are staying within the limit of 20 frames at constant resolution for the flow solver which converges in about 100 iterations. Using this framework we achieve an average computational time of one minute per-frame, roughly split into 20sec per flow solver and 40sec post-processing on a workstation with 16GB RAM 3.73 GHz CPU. We note that these times are an order of magnitude faster than those reported by Wand et al. [2007] and about
twice as fast as those of the recent marker-based method of White et al. [2007]. The memory requirements of our method are up to 4.5GB for the flow-solver and up to 2GB for the post-processing.

To generate the synthetic inputs in our experiments, we use a virtual multi-camera that simulates a real-time laser scanner. For each camera it shoots rays, registering the closest visible surface points from the virtual camera position. We record only position information for each scan point and not surface orientation.

**3D×time results.** Figure 1 shows the virtual scan of a running man, using four cameras. The large holes due to occlusions are completed in a natural manner with the space-time model.

Figure 11 shows a scan of a wooden doll, this time using only two virtual cameras. The larger occluded regions are again completed in a manner consistent with other frames, although this time the smooth completions in occluded regions are somewhat noticeable. As demonstrated, using an implicit FEM reconstruction alone (at \(128^3\) resolution) produces significantly inferior results.

Figure 3 shows a reconstruction of a moving hand puppet, captured by a low resolution structured light scanner from only two view points. The point data is noisy and has a persistent wide strip of missing data on the sides of the head. The front-facing arm sometimes occludes the face, and the back arm is sometimes occluded. The topology of the head and both arms is correctly reconstructed with the flow solution, and the geometry is fairly consistent. The thin palms of the puppet are too small to be recovered well given the flow grid resolution.

We also tested our method on inputs for which our incompressibility assumption is only approximately satisfied, such as the garment scans in Figures 12 and 13. Leveraging our iterative solver’s ability to solve the system inexactly, even in the case of singularities, our method succeeds in reconstructing these inputs and filling holes due to occlusion. In Figure 12, we use an animation of a jumping pair of pants [White et al. 2007] to create a simulated scan using three cameras. The pants rotate, so that almost every part of their surface is visible at some time. The flow solution accurately integrates the data from different times.

Figure 13 shows the results of applying our method to real scans of moving garments. The data was captured using a set of sixteen high-definition video cameras combined with a multiview stereo registration step [Bradley et al. 2008]. Although the scans contain holes due to occlusions, irregular sampling and noise, our flow system successfully generates a coherent 4D volume that smoothly varies across time and fills the missing data. Furthermore, due to the volumetric nature of our solution, the reconstructed shapes are always watertight manifold polygonal surfaces.

Finally, in Figure 10 we graph the mass convergence over the course of the flow iterations. The large jump in the first iteration arises from our initialization procedure, which marks cells known to be either inside or lying on the boundary as having a mass of 1. After the first iteration, we have a better guess of the amount of mass actually contained in the system. Therefore, as the iterations progress, the total amount of mass levels out. Similarly, Figure 4 shows the mass propagation from an initial guess to the actual mass in the last iteration. The measurements were carried on the Elephant data set [Wand et al. 2007] shown in the accompanying video. We note that while on this example both methods correctly reconstruct the watertight surface, an important feature of our method is that it always guarantees a watertight output which is not the case for Wand et al. [2007].

**Discussion and Limitations** We observed above that volume is not perfectly preserved by our reconstruction. There are several reasons for this. The bounded speed assumption (Section 3), which might appear to be strictly enforced by our formulation, is not perfectly satisfied by several of the inputs we generated, with data moving more than one-cell distance between frames. Also, for computational reasons and to guarantee a valid characteristic
function, we clamp material values to zero and one, potentially violating the incompressibility constraints in the process. Finally, as described in Section 5, the convergence tolerance, chosen to provide a sensible balance between accuracy and computational cost, is significantly larger than the machine roundoff error.

While these considerations do mean that our solution does not precisely satisfy the assumptions of the model, this flexibility in fact proves to be extremely useful. We can handle inputs that do not strictly meet the speed assumption or for which the volume does vary. Thus, when the constraints cannot be satisfied exactly, they are satisfied in a least squares sense. Nevertheless, in all our tests, the iterative solver reached a tolerance that maintains several decimal digits of accuracy. Indeed, in all our experiments we have found that the mass conservation error, measured as the square root of mass variance across the frames, was always less than 7% for the 2D examples and less than 2% for the 3D examples. The worst example in 2D is shown in Figure 7, in which the object does indeed experience a small change in mass when it splits.

While our post-processing step produces a generally smooth 4D solid, the final 3D meshes representing the time-slices of the solid are computed independently. Thus the connectivity can change between consecutive frames. In low resolution, this can cause some visual flickering. However, as the volume is refined, this flickering becomes less noticeable, and it eventually disappears in high resolution. Algorithms for temporally coherent meshing are an interesting research direction.

### 7 Conclusions

We have presented a novel volumetric approach for space-time surface reconstruction that leverages the knowledge of object behavior across time, to plausibly reconstruct the deforming surface despite persistent occlusions. Based on reasonable physical assumptions we formulate the reconstruction problem as a solution to a space-time formulation linking together mass and flow across the entire data sequence. We provide an efficient mechanism to solve the resulting optimization problem and demonstrate the viability of our approach on a set of complex examples. We compare our reconstruction results to those generated by extending two 3D reconstruction techniques to 2D x time (RBF and FEM) and 3D x time (FEM). A key idea in our work is to handle the time dimension very differently than the spatial dimensions using several flow priors. Hence, our method correctly completes gapping holes in the data which persist across multiple frames.

Despite the use of a specifically tailored solver, our method is still not suitable for tackling very large scale problems. One avenue for future work which will make it possible to solve problems with much higher resolution would be to consider a hierarchical or domain decomposition approach for optimizing the flow formulation.

Specific applications such as modeling moving humans [Carranza et al. 2003; de Aguiar et al. 2007] could be enhanced by incorporating additional assumptions, such as skeletal structure or rigidity, within the framework of material flow.

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### References


### Table 1: Number of frames, average number of boundary points per frame (ppf), and number of cells initialized over the whole volume for the 3D × time data sets presented.

<table>
<thead>
<tr>
<th>Data set</th>
<th>Frames</th>
<th>Avg. ppf</th>
<th>initialized</th>
</tr>
</thead>
<tbody>
<tr>
<td>Running man</td>
<td>30</td>
<td>21600</td>
<td>24.8%</td>
</tr>
<tr>
<td>Wooden doll</td>
<td>30</td>
<td>8000</td>
<td>20.7%</td>
</tr>
<tr>
<td>Pants</td>
<td>280</td>
<td>10100</td>
<td>29.8%</td>
</tr>
<tr>
<td>Puppet</td>
<td>200</td>
<td>34000</td>
<td>28.5%</td>
</tr>
<tr>
<td>Dress</td>
<td>130</td>
<td>31000</td>
<td>30.1%</td>
</tr>
<tr>
<td>T-shirt</td>
<td>160</td>
<td>35000</td>
<td>27.1%</td>
</tr>
<tr>
<td>Sweater</td>
<td>70</td>
<td>48000</td>
<td>27.3%</td>
</tr>
<tr>
<td>Elephant</td>
<td>20</td>
<td>40000</td>
<td>22.4%</td>
</tr>
</tbody>
</table>

Running man
Wooden doll
Pants
Puppet
Dress
T-shirt
Sweater
Elephant
Figure 11: Space-time surface of a 3D wooden doll’s walking cycle. Each row represents a particular frame from the cycle. (a) Points captured by the scanner. (b) The result of a basic FEM reconstruction. (c) Initial “inside” cell labeling. (d-f) Three solver iterations showcasing mass concentration within the surface as the solver progresses. (g) Final reconstruction.

Figure 12: Reconstruction of jumping pants from a motion capture sequence. Top: The points used for the reconstruction. Center: The mass field obtained at the final iteration of our flow solver. Bottom: our reconstruction.

Figure 13: Excerpts from reconstruction sequences of various moving garments captured from multiple video cameras. Left: the captured points. We show zoom-ins of irregular sampling, holes and outliers on the leftmost column. Right: our reconstruction.


Hansen, P. C. 1998. Rank-Deficient and Discrete Ill-Posed Problems. SIAM.


