IEEE Visualization 2004 Tutorial 4

Advanced Virtual Medicine: Techniques and Applications for Medicine-oriented Visualization

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Tutorial Summary

One of the largest application domains of visualization is medicine. Numerous techniques ranging from medical imaging to virtual medicine are used in both daily health care practice and in research. In particular, recent developments in minimally-invasive surgery require an advanced planning and intra-operative support through computer science methods. Even until recently, 2D medical imaging remained to be the standard for the planning of these interventions, while 3D and virtual medicine methods are only slowly merging into hospitals. This however, has dramatically changed with the introduction of intra-operative navigation systems.

In this course, we will first give an introduction into medical imaging methods such as data acquisition, data analysis, segmentation, registration and rendering both in 2D and 3D. Based on this foundation, the course will further explore a variety of applications for virtual medicine. In particular, we will discuss virtual endoscopy, virtual reality methods for surgery, soft-tissue-, and motion simulation, functional Magnetic-Resonance Imaging some of the most actively researched fields in virtual medicine. Together, these topics form important components towards more realistic interaction with the virtualized human body. Besides the technical aspects, we will also discuss the advantages to traditional methods, but will also illustrate their specific and inherent limitations.

For updates and additional information, see http://www.gris.uni-tuebingen.de/~bartz/tutorials/vis2004

1 Prerequisites

We assume a basic understanding of 3D graphics, visualization and medical-imaging technology. Although recent research results will be presented, we are not in particular targeting at experts in the field.

2 Tutorial Speaker

The material of this tutorial has been prepared by the following six people:

Dirk Bartz is head of the research group on Visual Computing for Medicine (VCM) of the University of Tübingen. His recent works covers interactive virtual medicine and scientific visualization. In 2002, he received the NDI Young Investigator Award for his work in virtual endoscopy and intra-operative navigation.

Dirk studied computer science and medicine at the University of Erlangen-Nürnberg and the SUNY at Stony Brook. He received a Diploma (M.S.) in computer science from the University of Erlangen-Nürnberg, and a Ph.D. in computer science from the University of Tübingen (both in Germany).

Jan Hardenbergh has worked in the 3D computer graphics industry for 25 years. He now works on software for VolumePro hardware engine at TeraRecon, Inc. He started when the project was still in research at MERL. He also helped bring up the Apollo DN10KVS and Oki TrianGL chips, both performance leaders for brief times. He has worked on the integration of 3D and the Internet by implementing products and serving on the architecture boards of both VRML and PEX (PHIGS and X). Jan also has a rich CAD (Design) background working with Computervision, Cognition and Revit and looks forward to working in CAD (Diagnostics).

Michael Hauth is a member of the research staff of the Computer Graphics Laboratory (GRIS) at the University of Tübingen. His current research focuses on simulation of deformable objects with applications in textile and medical animation. Michael studied computer sciences, physics and mathematics at the University of Tübingen. He received a Diploma in computer sciences in 1999 and a Diploma in mathematics in 2000. In 2000/2001 he held a visiting post as research assistant at the Numerical Analysis Group of the University of Geneva. He authored and co-authored several papers covering simulation and animation of deformable objects.

Klaus Mueller is currently an Assistant Professor at the Computer Science Department at Stony Brook University. His current research interests are computer graphics, medical-imaging, and virtual and augmented reality. He has authored or co-authored more than 30 conference and journal papers on volume rendering and on filter design. One of these papers won the Visualization "Best Paper" award in 1997. He has also won the Visualization "Best Hot Topic" award in 1999 and the NSF CAREER award in 2000. He has cochaired the 2001 and 2003 Volume Graphics Workshop and the 2002 Symposium on Volume Visualization and Graphics.

Markus Wacker is a member of the research staff of the Computer Graphics Laboratory (GRIS) at the University of Tübingen and currently head of the Virtual TryOn project in Tübingen. His research focuses on physically correct simulation of deformable objects for textile and medical applications. Markus studied mathematics and physics at the University of Tübingen and received a Ph.D. in mathematics from the University of Tübingen.

Yin Wu is a principal computer scientist in Terarecon Corporation, which is an imaging technology company that develops professional imaging applications, with special emphasis on medical imaging. Prior to her work at Terarecon she works on volume rendering and medical visualization in Mitsubishi Electric Research Lab (MERL). She received her Ph.D in Computer Science at University Geneva at 1998 on facial animation and biomechanics based aging simulation. Her research and development focus on physical simulation and interactive medical imaging visualization.

3 Schedule

The presentation itself is organized in four parts:

I.	Welcome and Introduction (Bartz, Wu)	60min
	Examples of Virtual Endoscopy Applications	
	Time Density Analysis of Tumors, Cardiac Time Volume Analysis, fMRI	
II.	Foundations	120min
	Medical Imaging Techniques (Bartz, Wu)	20min
	(X-Rays, Computed-Tomography, (functional) Magnetic-Resonance	
	Imaging, Rotational Angiography, Sampling-Theory)	
	Data Pre-processing (Bartz)	60min
	(Segmentation, Classification, Registration, and Fusion)	
	Coffee Break	
	Visualization and Navigation Techniques (Mueller, Wu)	40min
	(Indirect and direct surface volume rendering,	
	camera path generation vs. free fly-throughs	
	Lunch Break	
III.	Applications of Virtual Medicine	
	Virtual Endoscopy (Bartz)	20min
	(Available Systems, Advantages and Limitations of Virtual endoscopy)	
	Simulated Surgical Procedures (Wu)	40min
	fMRI for Surgical Planning (Hardenbergh)	45min
	Coffee Break	
	Soft-Tissue-Simulation (Wacker, Hauth)	90min
	General Basics	15min
	Physical and mathematical Background	45min
	State-of-the-art in simulation of Human Soft-Tissue	20min
	Soft-Tissue for minimally-invasive Surgery Simulation	10min
IV.	Questions and Answers (All)	15min

4 Introduction

These notes introduce into some topics covered in this tutorial. In particular, it introduces concepts of volumetric data representation and volume rendering (Section 5). Thereafter, Section 6 will introduce various scanning modalities. An overview of the various virtual medicine applications, in particular focusing on the skull, is given in Section 7. For a detailed overview on virtual endoscopy, we refer to the respective Eurographics 2003 STAR-report, which is included in the additional material on the conference DVD. Furthermore, the appendix contains interesting WWW links on the topic, to related papers, and – of course – the slides of the presentation. However, the webpage mentioned above probably contains a more up-to-date version of the presentation.

5 Volume Data

• Dirk Bartz, University of Tübingen, Germany

Traditionally, computer graphics represented a model as a set of vectors which were displayed on vector graphic displays [Foley et al. 1996]. With the introduction of raster displays, polygons became the basic rendering primitive, where the polygons of a model were rasterized into pixels, which represent the compounds of the framebuffer. The concept of an image composed of two-dimensional discrete pixels was later extended into a volume, composed of discrete voxels, arranged in a three-dimensional array. The representation of model data as a volume is particularly useful for scientific visualization, since the data is usually computed by a simulation on a multi-dimensional grid with three or more dimensions, or measured by a volumetric scanner like a CT scanner.

In the following sections, we will briefly introduce into the concepts of volume datasets (Section 5.1) and volume rendering. Specifically, we describe the two primary approaches to generate a visual representation of the volume dataset; indirect (Section 5.3) and direct volume rendering (Section 5.4). Both approaches use voxel-based shading, which will be reviewed in Section 5.2.

5.1 Volume Data

A volumetric dataset is a three-dimensional array of data values, the voxels. We address these voxels with three indices i, j, k, indicating the position of a voxel within the volume dataset. If we assume an array of 512 data values in each dimension, each index runs from



Figure 1: Volume cell in a rectilinear grid

i, j, k = 0..511, forming a $512 \times 512 \times 512$ volume dataset. The three-dimensional array can also be seen as a stack of two-dimensional arrays of data values and each of these two-dimensional arrays as an image (or slice), where each of the data values represents a pixel. This alternative view is motivated by the slice oriented, traditional way physicians look at a volumetric dataset).

Eight neighboring voxels $(V_{i,j,k}, V_{i+1,j,k}, V_{i,j+1,k}, V_{i+1,j+1,k}, V_{i,j,k+1},$

 $V_{i+1,j,k+1}$, $V_{i,j+1,k+1}$, $V_{i+1,j+1,k+1}$) form a volume cell, or simply cell¹ (see Fig. 1). The pixel distance between the voxels within one slice or image is usually equal. However, the slice distance between two neighboring voxels in two neighboring slices is frequently not equal to the pixel distance. We call these datasets anisotropic volume datasets. If slice and pixel distance are equal, we speak of an *isotropic* volume dataset. The distances are also called voxel spacing. In most datasets, all voxels are aligned on a *cartesian grid*, according to voxel spacing. Therefore, we speak of a cartesian or rectilinear grid dataset. If the voxel spacing is constant, the grid is also called an uniform grid, or a non-uniform grid vice-versa. Some scanners (i.e., 3D ultrasound) produce volume datasets which are not aligned on a cartesian grid, but on a grid which is bent. In other words, the grid geometry has changed, but the grid topology (connectivity) has remained the same. These grids are called curvilinear grids. Together with cartesian grids, they are classified as regular grids. If also the topology is no longer of a cube-like (possibly bent) cell, the resulting grids are classified as irregular or structured grids. If the grid topology is using various cell types of different connectivity, we speak of an unstructured grid. In the course of this thesis, we will only encounter regular, cartesian gridded volume datasets.

5.2 Voxel-based Shading

The shading of an object has a significant influence on the appearance of the object, since it simulates the interaction of light and the material properties of the object. Besides material and light, the shading is determined by the normals of the object, which define its surface. In traditional computer graphics, these normals are defined at the vertices of a triangle and they are interpolated over the area of the triangle to compute the shading of the triangle.

In contrast, the normals of a volumetric representation are defined at the positions of the data values. Due to the lack of a specific surface in a volume dataset, the normals are approximated by gradient operators, which describe changes in the material properties of a volume dataset. This method was originally motivated by the *partial volume effect* (see Appendix 6), which generates smooth density changes between two different materials (i.e., bone and tissue) in CT datasets [Höhne and Bernstein 1986], thus the density gradient represents the surface of the material interface between bone and tissue. The partial volume effect also affects other image modalities, such as MRI, or rotational angiography (see Appendix 6).

The currently used standard gradient operator – which we also use later on – computes the central difference between the six direct neighbors of a voxel [Höhne and Bernstein 1986], without taking into account the value of the voxel itself. Other gradient operators include the intermediate difference operator (sometimes also called forward or backward difference operator) – which consider the voxel values at the current position and at three neighboring voxels [Lichtenbelt et al. 1998] – and the Sorbel operator, which considers the full 26-neighborhood of a voxel to approximate the gradient [Lichtenbelt et al. 1998].

Once the normals at the voxel positions are computed by the (normalized) gradient operators, the normals at the sample points within a volume cell are usually computed by trilinear interpolation of the normals at the eight voxels of this cell.

5.3 Indirect Volume Rendering

In indirect volume rendering (IVR), a "traditional" computer graphics representation is generated, usually by extracting an isosurface or an isocontour from the dataset. The respective graphics primitives (i.e., polygons) are then rendered using standard computer graphics hardware. The most popular approach of IVR is probably the Marching Cubes algorithm [Lorensen and Cline 1987], where each volume cell of eight

¹An alternative, somewhat outdated terminology describes a voxel as a cubic cell around a data value which extents half way to the next neighboring voxels. The basic difference is the notion of interpolation between the data values, which is nearest-neighbor (non-continuous) for the older scheme, and trilinear (continuous, or C^0) for the current scheme.



Figure 2: Reduced Marching Cubes case table [Kaufman et al. 1996]

neighboring voxels is classified according to the specified isovalue (which specifies the isosurface). 2^8 possible combinations whether the scalar value of a voxel of the cell is above or below (or equal to) the isovalue are stored in a table of 256 cases. In the original paper on Marching Cubes, these 256 cases were reduced to 15 cases by inverting or rotating the classification cubes (see Fig. 2). However, later research showed that these reduced case table generates inconsistencies, which result in holes in the isosurface [Kaufman et al. 1996]. Therefore, we only use a full 256 case table, similar to many other state-of-the-art implementations of the Marching Cubes algorithms [Schroeder et al. 1998].

Up to five triangles per cell are generated depending on the classification case. To approximate the normals at the vertices of the generated triangles, each normal is computed by a trilinear interpolation of the normals at the eight voxels of the current cell, which in turn are computed by a central difference operator (see Section 5.2) [Höhne and Bernstein 1986].

Several approaches addressed improvement of the basic Marching Cubes algorithm. Most of them focused on skipping of regions which do not contain the isosurface, by applying hierarchical data-structures, like quadtrees and octrees [Samet 1994], k-D-trees [Bentley 1975], and BSP-trees [Fuchs et al. 1980]. Wilhelms and van Gelder introduced the Branch-ON-demand-Octree (BONO) [Wilhelms and van Gelder 1992], which stores the isovalue interval of the octants to skip octants whose isosurface interval does not contain the specified isovalue.

Many other approaches are known for the efficient isosurface or contour extraction of large volumetric scalar fields. Value partitioning methods [Livnat et al. 1996; Shen et al. 1996; Cignoni et al. 1997] store the minimum and maximum values of the voxels of a cell as a pair in a 2D field. Livnat et al. used a k-D-tree structure [Livnat et al. 1996], and Shen et al. [Shen et al. 1996] used lattice subdivision to subdivide this field. Cignoni et al. used an interval tree as search index for optimal efficiency [Cignoni et al. 1997]. Space and value partitioning methods are used for an efficient contour propagation, starting from a small set of seed cells [Itoh and Koyamada 1995; Bajaj et al. 1996].

5.4 Direct Volume Rendering



Figure 3: (Post-shading) Direct Volume Rendering Pipeline

In contrast to IVR, methods of direct volume rendering (DVR) are generating images without an intermediate polygonal representation.

Instead, the volume dataset is projected onto an image- or view-plane. Figure 3 shows an overview of the direct volume rendering pipeline using a post-shading scheme. After computing a sample according to the chosen algorithms, it is *classified* by the transfer functions and shaded according to the specified lights. Finally, the lit sample is *composited* with the previous samples using the over-operator [Porter and Duff 1984].

The transfer functions define the contribution of the samples by mapping their value(s) to a red, green, blue, and α (transparency) value. By evaluating the volume rendering integral (see Equation 1), the contribution of the volume data (through the transfer functions) lit by the possibly multiple light sources is computed for each pixel on the image- or view-plane, where I_{λ} is the color intensity of the pixel at \vec{x} , which receives a contribution of a ray of the length L – casted from \vec{x} in direction \vec{r} – and λ corresponds to the wavelength of a color (r, g, b). $C_{\lambda}(s)$ is the actual color of the sample at position s on the ray, with an extinction coefficient $\mu(s)$ at the sample, which is basically the opacity of the sample. The opacity-weight color is attenuated by the exponential opacity term, which collects the opacity of the ray through the volume (from entry point 0 to current sample s).

$$I_{\lambda}(\vec{x},\vec{r}) = \int_{0}^{L} C_{\lambda}(s)\mu(s)e^{\left(-\int_{0}^{s}\mu(t)dt\right)} ds$$

$$\tag{1}$$

Unfortunately, the analytical volume rendering integral cannot be computed for the general case [Max 1995], hence a discrete approximation is used, where Δs is the width of the discretized integration interval:

$$I_{\lambda}(\vec{x}, \vec{r}) = \sum_{i=0}^{L/\Delta s} C_{\lambda}(s_i) \mu(s_i) \prod_{j=0}^{i-1} e^{(-\mu(s_j)\Delta s)}$$
(2)

The theoretical basis of the volume rendering integral is the *density emitter model* introduced by Sabella [Sabella 1988], which assumes a simplified model of the transport theory of light [Hege et al. 1993]. This simplified models takes only absorption and emission into account; physical terms like scattering, influence of the different wavelengths (red, green, and blue), or the influence of participating media are ignored.

Generally, DVR-algorithms are classified into front- and back-projection approaches. While front-projection approaches project the contributions of a dataset onto the view-plane, back-projection approaches trace the contribution for each pixel of the view-plane through the volume. Prominent candidates for front-projection are splatting [Westover 1990; Westover 1991; Mueller and Crawfis 1998], texture-mapping [Cullip and Neumann 1993; Cabral et al. 1994], or cell projection [Wilhelms and van Geldern 1991]. For back-projection, the most prominent candidate is ray-casting [Tuy and Tuy 1984; Levoy 1988].

Another important algorithmic classification is the shading scheme, which basically depends on when and how the normals at the sample points are calculated. Most approaches use the normalized gradient of the sample, based on central differences, to compute the normal at the sample [Höhne and Bernstein 1986] (also see previous section on voxel-based shading). However, this gradient can be computed before (pre-shading), or after the sample is computed (post-shading), as described above. Pre-shading pre-computes the normal gradients – and hence the illumination and shading – for all voxel values of the volume. The resulting colors and opacities are stored as a color volume instead of the original data. During the actual rendering process, the samples are computed based on the color volume. If the lighting conditions are changing (i.e., modifications of the lights for a diffuse lighting model, or of the view-point in a Phong-like lighting model), this requires the re-calculation of this color volume. Additionally, the colors need to be pre-multiplied by the opacities in order to avoid color bleeding artifacts [Wittenbrink et al. 1998] (see Fig. 4). In contrast, post-shading computes the classification and normals during rendering, after computing the sample.



Figure 4: Color bleeding artifacts in a rotational angiography dataset. (a) Non-opacity weight, pre-shaded volume rendering; color bleeding artifacts are marked with red ellipse. (b) Post-shaded volume rendering.

In the following, we will briefly describe four of the most popular direct volume rendering approaches, where each of them has their own specific advantages and disadvantages.

Ray Casting



Figure 5: Ray casting of a volume dataset with parallel projection and uniform sampling.

In the image-space oriented ray casting approaches, rays are cast from the view-point through the view-plane into the volume. Along their way through that volume, samples are calculated usually at equal sampling distances between two sample points (see Fig. 5). A sample is computed based on trilinear interpolation within a cell of eight voxels. Thereafter, it is classified according to the transfer functions. If that sample has a contribution to the ray, the normal gradient is computed based on a trilinear interpolation of the normalized central differences at the eight voxels of the cell which contains the sample point. Finally, the sample is composited with the previous samples of the ray.

Acceleration methods include *early ray termination*, where the sampling along the ray is terminated, once (almost) full opacity has been reached, or space leaping, where a distance field or other data-structures indicate empty space where no sampling is required. Ray casting can also utilize oversampling within the view-plane and along the ray to account for high-frequency data in the data volume.

Shear-Warp Factorization



Figure 6: Shear-warp Factorization: (a) The volume slices are shifted (sheared) to account for the angle between view- and base-plane. (b) The base-plane image is warped to the view-plane.

A variation of the ray casting approach is the image-space oriented shear-warp factorization, which essentially factorizes the viewing transformation into a shearing and a warping matrix [Lacroute and Levoy 1994]. This enables the casting of rays from each pixel of a volume aligned base-plane, which is most parallel to the image- or view-plane, which in turn can be exploited for an optimized memory access. To take into account the angle between view- and base-plane, the individual slices are shifted (**sheared**) accordingly (see Fig. 6a). The samples of a whole scanline are computed simultaneously within one slice, employing a bilinear interpolation scheme. This results in a view-dependent sampling interval, which can vary between 1.0 for axis-aligned views, to $1.73 (= \sqrt{3})$ for corner-on views. Perspective projection also requires the scaling of the slices to address the divergence of perspectively cast rays. Finally, the computed base-plane image is **warped** onto the view-plane (see Fig. 6b).

The Stanford VolPack-implementation of the shear-warp factorization is highly memory optimized. A pre-computed classification and a run-length-encoding of the opacity weighted voxels are used to rapidly skip the transparent, non-contributing volume space. VolPack uses a pre-shading scheme which computes the lighting on the pre-classified voxels, before generating the sample within a slice. Note that this (pre-)shading operation occurs during rendering, not in a pre-process, which allows the modification of the light settings without re-generating the pre-classified volume dataset.

Splatting



Figure 7: Splats arranged on view-plane aligned voxel slaps are projected into sheet buffers, which are in turn composited front to back into the final image on the view-plane.

The object-space oriented splatting approach projects each voxel onto the screen as an overlapping, orientation invariant Gaussian kernel with an amplitude scaled according to the voxel value [Westover 1990; Westover 1991]. High speed is obtained by a footprint lookup table of the pre-computed, radially symmetric kernel function. For correct compositing, the volume is processed by slices oriented most parallel to the view-plane, which causes severe brightness variations, such as popping artifacts in animated views. Mueller et al. modified the splatting approach such that *slabs* of the voxel kernels were processed in an image aligned fashion [Mueller and Crawfis 1998] and projected in *sheet buffers*, which in turn are composited into the final image on the view-plane. Other modifications include the *early splat elimination* for the removal of non-contributing splats from rasterization [Mueller et al. 1999]. Similar to early ray termination, the (average) opacity of the screen area covered by a splat is tested, if it is above a specified threshold. In this case, the respective splat will not have a significant contribution (or no contribution, if the chosen threshold specifies full opacity). The test itself is performed by convolving the associated opacity buffer area (which stores the up to now accumulated opacity of each sheet buffer) with a box filter of the screen size of the splat, resulting in the opacity average of that area [Mueller et al. 1999].

Instead of tri-/bilinear interpolation-based point sampling, splatting employs a sample average across the sampling distance in view direction. This introduces an additional low-pass filter operation, which reduces aliasing, but also tends to smooth signal characteristics. Finally, the recent splatting approaches [Mueller and Crawfis 1998; Mueller et al. 1999] also provide post-shaded DVR, in contrast to pre-shading provided by the original splatting approach.

Texture Mapping



Figure 8: 3D Texture Mapping-based Volume Rendering [Westermann and Ertl 1998]: (a) 3D texture slices are generated from the volume, perpendicular to the view-plane; the texture slices are mapped onto the screen (b) and blended with the previous slice (c).

Object-space oriented texture-mapping is accumulating texture slices perpendicular to the view-plane back-to-front using the blending functionality of graphics hardware (see Fig. 8) [Cullip and Neumann 1993; Cabral et al. 1994]. The sampling of the texture slices from the volume are either trilinear, if 3D texture mapping hardware is available, or bilinear, if only 2D textures are supported. (On some graphics systems, quadlinear interpolation is also available, where an additional interpolation takes place between two mipmap-levels.)

Shading can be achieved by the computation of a pre-shaded color volume [van Gelder and Kim 1996], or by using multi-pass methods to visualize isosurfaces [Westermann and Ertl 1998; Sommer et al. 1998; Dachille et al. 1008], or transparent volumes [Meißner et al. 1999]. In

our comparison later-on, we use the color volume approach to provide high-speed volume rendering of semi-transparent volumes. However, due to the limited precision of the alpha channel of the framebuffer (only eight bits), we do not pre-multiply the voxels with the opacity which leads to color bleeding artifacts [Wittenbrink et al. 1998] (see Fig. 4).

6 Volume Data Acquisition Techniques

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There are many sources of data for medical visualization. For our purposes, we rely on data from medical scanners, such as Computed Tomography (CT or CAT scanners), Magnetic Resonance Tomography (MRT or MR scanner), rotational biplane X-ray (or rotational angiography), 3D Ultrasound, Positive Emission Tomography (PET), and many more. These scanners produce a stack of images, where each image represents a slice from a data volume. Each entity or pixel of the images represents a sample point within the data volume and it is also called a voxel (see Section 5).

In the course of this section, we give an introduction into basic concepts of the scanning technology of the modalities of the volume datasets later used for VIVENDI. More details on the specific technology can be found in [Cho et al. 1993; Lehmann et al. 1997; Kalender 2000]. Several sources of artifacts are known; most notorious are aliasing problems due to undersampling, motion artifacts if the scanned object is moving during the scanning procedure (i.e., a heart), and *partial volume effects*, which are also related to undersampling of a structure. 3D data samples (voxels) which are reconstructed from projections of a 3D scanner are averages of the local volume. If material with a high voxel value is adjacent to material with a low voxel value, this averaging can lucidly distort these voxel values, causing the partial volume effect.

6.1 X-Ray

In classical X-ray imaging, electrons are shot at the focal spot of a specific target. The resulting radiation is absorbed (and scattered) by an object (i.e., body parts) behind the focal spot and hence the attenuated intensity is recorded on detectors behind the object. Depending on the quantity of the acceleration voltage, X-rays are denoted as hard or soft radiation. The interaction of X-rays with matter is basically determined by photo-electric absorption and by scattering (Compton scattering). With hard radiation, the scattering effect is dominating the interaction, resulting in an absorption only significant in object areas with high density (high atomic number), i.e. bones. For areas with a lower density (i.e., tissue), a sufficient absorption requires soft radiation. Today's X-ray devices feature a very high resolution (i.e., 4096×4096 pixels), but only as a 2D projection.

A variation of classical X-ray imaging is fluoroscopy. The emitted radiation is recorded with an image intensifier and displayed such that the progress of specific procedures can be observed. This permanent exposure to ionizing radiation requires a significant lower intensity than with classical X-rays. Hence, the resolution and general image quality are reduced.

6.2 Computed Tomography

The introduction of X-ray Computed Tomography (CT) in 1972 [Hounsfield 1972] provided for the first time a volumetric representation of objects, and not only a 2D projection of a volumetric object. Generally, it is seen as one of the major milestones in medical imaging [Cho et al. 1993].

As basic concept, X-rays are emitted through the object from different positions around that object (i.e., filling an angular range of 180 degree) and the intensity profile is recorded by a detector. Algorithms are applied to reconstruct an intensity attenuation layer of the object. A series of these projection with an incremental modified position generates a stack of intensity attenuation layers, which forms an intensity volume. The intensity attenuation is measured in Hounsfield units (HU), with water as reference material (0 HU). I.e., bone has a high measured intensity attenuation, while fat or air have negative HU [Lehmann et al. 1997]. Each of the reconstructed volumetric samples (voxels) represents the average of the local environment.

The major differences between the development stages of CT [Kalender 2000] are different projection reconstruction algorithms and the emitter and detector architecture (data acquisition). The first generation of CT scanners was basically the experimental setting of Hounsfield's CT scanner. It used a single pencil-like X-ray beam emitter and a single detector on the opposite side of the object. To acquire a data slice, the pencil beam is translated along the object and rotated afterwards for the next series of beams. All together, the costly mechanical movement of emitter and detector caused long scanning times, ranging from several minutes to several hours at a resolution of 80×80 pixels per scan/slice. Furthermore, the single emitter/detector architecture enabled only a poor utilization of the emitted radiation.

The next generation and first commercial generation of CT scanner used small angle fan beams and multiple detectors to scan two neighboring rotational projections at the same time. This technique reduces the number of necessary rotations, and hence the required scanning time (10 - 60 seconds, up to several minutes) needed for a sufficient reconstruction. It also provides a better utilization of the emitted radiation. Both first and second generation techniques are parallel beam devices, which use different reconstruction algorithms than the next fan beam devices.

The next improvement increased the fan beam angle and the number of detectors to cover the whole object, thus the translating movement became unnecessary and increased the scanning speed to five seconds per slice. Similar to the previous techniques, the radiation is enabled in fixed time intervals to be measured by the detectors.

In the fourth generation, the rotating detector was replaced by a fixed circular ring detector, which reduced the technical effort of moving the larger mass of emitter and detector. Here, the radiation was permanently emitted and only the detectors were enabled at certain intervals. However, several problems of ring detectors led to further developments in favor of third generation scanners. Besides the higher costs for the detector ring, specific X-ray scattering problems reduced the image quality of these systems, while collimator technology could reduce the scattering problems with a rotating emitter/detector system of the third generation.

Currently, the state-of-the-art are spiral or helical CT, where the emitter/detector system is rotating permanently around the object, while the object is moved continuously in the perpendicular direction to acquire a full data volume. This technique enables faster scanner due to the continuous rotating movement of the emitter/detector system which saves the time for the previously needed time to accelerate and slow down these heavy parts of the scanner. Recently, multiple layers of emitters and detectors (twin or quad slices) where combined to create multi-slice CT scanners, which enable fast and isotropic scanning of large object areas.

Besides the architectural development, different volume/slice reconstruction algorithms differentiate the various systems and generations. The basic approach is the back transformation of the slice projections into the volume slices by the Radon transform [Cho et al. 1993]. In the beginning of computed tomography, algebraic reconstruction techniques (ART) were used to solve this back projection problem. However, the high computational costs of iterative solving the large matrices – the size of a reconstruction matrix is equal to the resolution of the slice – rendered this approach as not usable for standard applications [Kalender 2000]. The standard method today is filtered back-projection, where each projection is composed according to the measured direction. Parallel and fan beam methods are available to perform this back-projection. However, the current fan beam methods are more complex and less efficient than state-of-the-art parallel beam reconstruction algorithms. Hence, the projections of today's fan-beam scanners are re-sorted in parallel beams before the actual reconstruction. Another modification is required to address the continuously moving object tray of modern spiral CT scanners, where a z-interpolation corrects the measured projections according to the tray movement [Kalender 2000]. In the future, cone beam reconstruction algorithms will probably replace the current methods [Kalender 2000], which are already successfully used in rotational biplane X-ray (see below).

6.3 Rotational Biplane X-Ray

Rotational biplane X-ray is a recent scanning technology which started from digital subtraction angiography (DSA – in DSA, X-ray images are recorded before (mask) and after (filling) the injection of a contrast agent. The mask is subtracted from the filling image, thus presenting only the contrast agent enhanced object areas). Hence, most of the associated applications are angiography applications and this technique is frequently referred to as rotational angiography [Fahrig 1999; Gürvit et al. 2000].

To generate volumetric datasets, a series of up to 132 X-ray projections are taken from a rotation range of 200 degrees around the scanning object. In contrast to CT, rotational biplane X-ray is using a full array of up to 1024^2 detectors which enable the measurement of a full cone of rays. To account for reconstruction errors [Kalender 2000], a modified back-projection algorithm is used [Hornegger 2000]. Additionally, special filter kernels are used which further reduces potential artifacts. Current rotational angiography systems provides very high resolution, isotropic datasets, good reconstruction quality, and a high data acquisition speed of up to 13 seconds for a full scan.

6.4 Magnetic Resonance Imaging

The beginning of Magnetic Resonance Imaging (MRI) dates back to the early seventies, but it was not adopted into medical use until the nineteen-eighties. It is based on the nuclear resonance of hydrogen in a magnetic field, where each of the hydrogen nuclei can be considered as a small dipole magnet which aligns itself either parallel or anti-parallel along the magnetic field.

While aligned in that field, the protons (which are identical to the hydrogen nucleus) spin arbitrarily around the axis of the field. This spinning is called the *precession* of the nuclei. If energy is applied to the magnetic field as a radio-frequency pulse (RF) at the *Larmor*-frequency, the nuclear resonance forces the protons to receive some of the energy from the RF. This pulse also forces all the protons to spin synchronously, or *in phase* and flip increasingly into the anti-parallel orientation of higher energy, until the number of parallel protons is equal to the number of anti-parallel oriented protons. The duration of the RF pulse determines the amount of precession; i.e., a *90 degree pulse* will force the protons into a 90 degree precession, where the precession vector of the protons, they slowly release the received energy, *de-phase* and re-align with the magnetic field. This *relaxation* is described as *free induction decay (FID)* and is divided into the transverse and longitudinal relaxation. The first relaxation of the transverse magnetization – also called spin-spin relaxation – describes the de-phasing of the x/y-component of the precession. The time required for this relaxation is called T_2 and is in the order of a few milliseconds. The longitudinal, or spin-lattice relaxation describes the re-alignment of the precession with the magnetic field, thus the restoring of the z-component. This relaxation time is called T_1 and is in the order of seconds. The actual measured volumetric information is the proton density σ which needs to be reconstructed at the specific voxels.

To reconstruct the spatial information of the measured signal, two additional gradient magnetic fields are applied. The first field selects the slice in z-direction of the volume, since only one layer of protons suffices the Larmor-frequency for the main and gradient magnetic fields. An additional gradient field in x-direction selects a y-slab. With a 2D Fourier reconstruction, the x/y-coordinate is encoded into that signal [Lehmann et al. 1997] by increasing de-phasing signal of the transverse relaxation along the x and y directions, which generate specific frequencies into the signal. Thus, the RF intensity (proton/spin intensity) is encoded into the intensity of the signal, while the position is encoded into the frequencies [Cho et al. 1993; Lehmann et al. 1997].

Different protocols describe sequences of various RF pulses and the actual measurement of the signal, where the time between the initial stimulation and the measurement is called *echo time* T_E and the time between two (initial) stimulation cycles is called *repetition time* T_R . By varying T_E and T_R , different weight data can be achieved; i.e., with short echo and repetition times, the proton density is mostly dominated by the T_1 relaxation time (T_1 weighted), with long T_E and T_R , the T_2 relaxation time is dominating the signal (T_2 weighted), or with a long T_R and a short T_E , the resulting data is neither T_1 nor T_2 and can be seen as "the pure (proton) density function" [Cho et al. 1993]. Scanning protocols such as Turbo Spin Echo (TSE) or Constructive Interference in Steady States (3D CISS) belong to these categories. A slightly different technique is used for angiography sequences, which focus on flow information. Specifically, the Time of Flight sequence (TOF) measures the spin saturation of the protons; the repeated stimulation of the protons in the data volume leads to a saturation of the signal of stationary samples. However, the particle flow in blood vessels are continuously introducing "fresh material" into the magnetic field which gives a strong signal. Typical MRI scans take between 2 and 25 minutes. This relatively long scanning time is dominated by the relaxation of the spins, not by the time required for the measurements.

Appendix: Virtual Endoscopy Resources

This text is also contained as html-file on the conference DVD.

Books

There are not many books on virtual endoscopy. If you are only interested in the clinical aspect, but not at all in the technical aspects, the book by Patrick Rogalla can be helpful:

P. Rogalla (ed.): Virtual Endoscopy and Related 3D Techniques. Springer Verlag 2001.

Other books (in particular books for medical students) sometimes briefly touch the subject, but don't go deeper into the subject. My thesis contains some related information, but the relevant parts are included in the tutorial notes or STAR report anyway.

Web Resources

In contrast, there are plenty of web pages related to the topic of virtual endoscopy. This list however, contains only a small subset of it. Try web search engines for more information on the web. Also on the conference DVD, we will add a (subjective) bibliography of related papers, mostly of our groups, with weblinks to the respective papers.

- http://www.gris.uni-tuebingen.de/ bartz/tutorials/vis2003: Course home page
- http://www.gris.uni-tuebingen.de/bartz/endoscopy.html: Virtual Endoscopy Project, University of Tübingen (VIVENDI Project)
- http://www.cs.sunysb.edu/ vislab/: Center for Visual Computing, SUNY Stony Brook (VICON Project)
- http://www.vec.wfubmc.edu: Virtual Endoscopy Center, University of Wake Forest (FreeFlight Project)
- http://3dradiology.org: Medical Imaging Lab, Stanford University
- http://splweb.bwh.harvard.edu:8000: Surgical Planing Lab, Brigham and Women's Hospital
- http://cobb.ee.psu.edu/users/krishnan/quicksee/quicksee.html: Multi-Dimensional Image Processing Lab, Pennsylvania State University
- http://www.rad.unipi.it/virtual/VE-forum/: Virtual Endoscopy Forum
- http://www.cg.tuwien.ac.at/research/vis/vismed/: TU Vienna/TIANI Virtual Endoscopy Projects

Medical Imaging Links

- http://www.kitware.com/vtk: Visualization Toolkit (VTK)
- http://www.itk.org: National Library of Medicine's Insight Segmentation and Registration Toolkit (ITK)
- http://www.image-registration.com: CISG registration of King's College, London
- http://dicom.offis.de/index.php.en: OFFIS-DICOM toolkit

Related Medical Imaging Companies

- http://www.viatronix.com: Viatronix
- http://www.vitalimages.com: Vital Images
- http://www.siemensmedical.com: Siemens Medical Solutions
- http://www.medical.philips.com: Philips Medical Systems
- http://www.gemedicalsystems.com: GE Medical Systems
- http://www.tiani.com: TIANI

Visit also the volume dataset repository at http://www.volvis.org www.volvis.org.

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Advanced Virtual Medicine: Techniques and Applications for Medicine-Oriented Visualization

Introduction

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Outline



Introduction (Bartz, Wu)

- Basics of Medical Imaging
 - Data Acquisition (Bartz, Hardenbergh)
 - Data Pre-Processing (Bartz)
 - Visualization and Navigation Techniques (Mueller, Wu)

Applications of Virtual Medicine (Bartz, Wu)

fMRI for Surgical Planning (Hardenbergh)

Soft-Tissue-Simulation and Deformation (Wacker, Hauth)

Questions and Answers (All)

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Additional Tutorial Notes

The conference DVD does contain additional material:

- Textual tutorial notes on data acquisition and volumetric datasets
- Virtual endoscopy state-of-the-art report of Eurographics 2003
- Links to related websites and papers (to avoid copyright clashes ...)
- and the slides, of course

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Virtual Endoscopy Applications:

- Colonoscopy
- Ventriculoscopy
- Angioscopy
- Bronchoscopy
-



















Virtual Ventriculoscopy (2)



Procedures:

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- placement of stent
 - requires replacement after a few years
 - causes degenerations of VS
- ventriculoscopy:
 - + no implantation required
 - complex procedure







Left Lateral Ventricle, approach from Posterior Horn: AH - Anterior Horn, CP - Choroid Plexus, CPV - Choroid Plexus Vein





Virtual Ventriculoscopy (5)





Foramen of Monro, approach from Third Ventricle: AI - Adhesio Interthalamica, LLV - entrance to Left Ventricle, RLV - entrance to Right Ventricle, LT - Lamina Terminalis



























Cardiac Time Volume Analysis



Cardiac Application:

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- Automatic contour detection of LV endocardial and epicardial surfaces
- Manual correction of the surfaces.
- Calculation of myocardial volume (and mass).
- Regional wall thickness, thickening, and ejection fraction can be displayed on polar maps and graphs.
- Time-volume graph, calculation of peak ejection, peak filling rate and their corresponding time.
- 3D display of endocardium and epicardium, with color coded regional indexes on the surface.

Cardiac Time Volume Analysis



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Advanced Virtual Medicine: Techniques and Applications for Medicine-Oriented Visualization

Foundations

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Foundations















Medical Imaging Techniques (8)



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Functional MRI (fMRI)

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- Blood flow increases to active regions of the brain saturates it with oxygen.
- \bullet Deoxyhemoglobin is paramagnetic (no $O_2)$ and can be imaged with fMRI.
- While in scanner, subject exercises mental functions.
- This is extremely useful in neurosurgical planning.



VIS04

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Functional MRI (fMRI)

- · Reconstructed scanning dataset includes volumes of
 - Anatomy of brain

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- Vasculature (blood vessels)
- Each volume for an activation area

 Medical Imaging Techniques (8)
 Many other imaging techniques:

 • (3D) Ultrasound
 • Positron Emission Tomography (PET)

 • SPECT
 •





























- Automatic segmentation frequently segments too much, or not all structures
- Manual segmentation is usually too expensive for daily practice (ie., visible human datasets)
- Semi-automatic segmentation with little interaction only: can consist of several steps

Check out: http://www.itk.org

Tutorial 4















Classification (1)

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- Classification specify how data is rendered (for direct volume rendering)
- Often confused with segmentation
- Are expressed by transfer functions
- Are usually based on **histogram**: every possible scalar value is assigned to a tuple of color and opacity
- Focuses on material interfaces
- May introduce high frequencies

Classification (2)
Example of transfer function
• Histogram:
• Rising or falling edges indicate material interfaces
• Gradient magnitude can be also provide useful information
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Data Fusion (2)

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Consider rendering parameters how to incorporate data from different sources:

- (Relatively) simple for surfaces
- Difficult for accumulative volume rendering

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Navigation and Rendering in Virtual Endoscopy

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Visualization Paradigms



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Available visualization paradigms

- 2D viewing, slice by slice (non-intuitive)
- 3D visualization (most appropriate)

3D visualization paradigms (use colon as example)

- Section colon into straight pieces, slice in the center, and visualize on a virtual tray
- Unroll and flatten colon and view as a 2D sheet
- Leave as is and perform a virtual fly-through

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 3D Fly-Through Paradigms
 Constrained of the second sec



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Potential Field Computation Merge the two potential fields Each voxel has potential according to both criteria • Pull user towards target and away from wall • Pull user towards target and away form wall • State of the sta

Rendering: Two Options



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Extract polygon mesh from boundary and visualize with graphics hardware

- Discards the original volume data
- Will not allow user to "drill" into the wall to reveal inside-structures
- Visualize the volume with direct volume rendering
 - Nothing is lost, "drilling" possible
 - Will also give more realistic (softer) images since there is no (linear) mesh approximation
 - Downside: computationally expensive









Rendering With Polygons: Algorithm With Occlusion Culling

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Locate cell containing camera, render its polygons

Initialize Aggregate Cull Rectangle (ACR) to screen





Rendering With Polygons: Algorithm With Occlusion Culling







Stop when ACR degenerates to zero









































Advanced Virtual Medicine: Techniques and Applications for Medicine-Oriented Visualization

Visualization and Navigation Techniques II Yin Wu

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Outline



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Introduction

Foundations

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Applications of Virtual Medicine

- Virtual Endoscopy
- Distributed Virtual Medicine
- fMRI for Surgical Planning
- Soft-Tissue Simulation /Deformation

Question and Answers

Techniques and Applications for Medicine-Oriented Visualization

Advanced Virtual Medicine:

Applications of Virtual Medicine Virtual Endoscopy Dirk Bartz

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Limitations of VE:

- 2. Rendering
- Rendering never better than data
- Resolution of real biopsy is unmatched
- Transparent rendering of inner surface ("look through", "virtual biopsy") does not replace real biopsy

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Advanced Virtual Medicine: Techniques and Applications for Medicine-Oriented Visualization

Applications of Virtual Medicine: Distributed Virtual Medicine

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Advanced Virtual Medicine: Techniques and Applications for Medicine-Oriented Visualization

Applications of Virtual Medicine: fMRI for Surgical Planning

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fMRI Scanning

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Done in a standard MRI scanner

Scan takes up to 2 hours to scan anatomy, vasculature and functional areas.

Patient performs various tasks, moving hands, reading, verbalizing.



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fMRI volume data Packing Activation data 8 channels is hard (especially with masking) 2 volumes + one for each functional area Example has three functional areas 40 MB of texture memory is getting smaller, but, still hard to fit in 64MB. · Right hand motor Language acquisition Pack the 3 activation datasets into 1. Verbal Generation Offset the voxel values (remember how) Each volume is 256x256x124 = 8MB. RHM 2 4 9 AC 14 16 5 VG 22 24 9 -VG-AC 36 36 4 **IEEE Visualization 2004** Tutorial 4 IEEE Visualization 2004 Tutorial 4















fMRI UI

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Alternatively, We can display the activation areas as independent Activations.

2			×	
	fMRI Control	Panel		
Vol	Functional MRI st ime must be creater	1 ort cuts 1 by Makel	MRI	
	see silver-quicks	start.pdf		
	Activation Visibility		1	
	🖙 Gray Matter (SPG	2		
	Vascular (VEINS2)	_	1	
	RHM	_	1	
	I⊽ AC	_	1	
	I⊽ VG	_		
	₩ -VG-AC	_	1	
	E Wardson Laurel Weise			
	- WINDOW Developen	•		
	Clip Plane			
	Mouse Moves Clipp	er		
	Save Image	,		









More Info

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This information as of August 2004: www.jch.com/volumes/fmriviz.htm

New books on fMRI published



Advanced Virtual Medicine: Techniques and Applications for Medicine-Oriented Visualization

Soft-Tissue-Simulation and Deformation

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 Current Activities
 Measurement

 • Enhance database
 • Comparative measurements [Ottensmeyer 2003]

 • Define standard benchmarks:
 Truthcube.org, E. Mazza

 • Define standard benchmarks:
 Truthcube.org, E. Mazza

 • Develop better in vivo instruments
 • Develop better in vivo instruments

 • Develop better in vivo instruments
 • Utorial 4

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Finite Differences / Finite Element Method



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- Toolbox to cast a physical formulation into a simulation
- Sound derivation from continuum mechanics
- Standard method in engineering
- Computationally expensive











Summary

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Finite Differences:

- Simple and easily optimized
- Moderate numerical approximation quality
- Difficult to fit to arbitray geometry
- Problem: Unstructured meshes

Finite Elements:

- Good numerical approximation quality
- For linear shape functions: $\varepsilon(u)$ =const per tetrahedron!

Good shape fitting properties
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Some Simulators	20 VIS 04 Rustin, toxes
State of the art: - Mist - Kismet / Surgical Science / Xitact	Commercial
- INRIA Epidaure	Research
- Lasso / CoMe	
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mages

INRIA/Epidau

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Lasso / CoMe
Two large Swiss National projects (Lasso 1996-2000, CoMe 2001-)
Soft tissue part focuses on hysteroscopy
Includes measurement subproject
Uses partly finite elements but also investigates customized mass spring models
Includes automated texture and pathology generation (e.g. fibroids), and special effects (e.g. bleeding)
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Directions of further work



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- (Better in vivo measurement devices)
- (Better/more complex force feedback devices)
- Cutting
- Adaptive Subdivision/Coarsening
- Numerical Methods (FEM, time integrators)
- Evaluation of haptic perception
- Parallelization





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