# Hardware-Accelerated Volume Rendering for Real-Time Medical Data Visualization

Rui Shen and Pierre Boulanger

Department of Computing Science University of Alberta Edmonton, Alberta, Canada T6G 2E8 {rshen,pierreb}@cs.ualberta.ca

Abstract. Volumetric data rendering has become an important tool in various medical procedures as it allows the unbiased visualization of fine details of volumetric medical data (CT, MRI, fMRI). However, due to the large amount of computation involved, the rendering time increases dramatically as the size of the data set grows. This paper presents several acceleration techniques of volume rendering using general-purpose GPU. Some techniques enhance the rendering speed of software ray casting based on voxels' opacity information, while the others improve traditional hardware-accelerated object-order volume rendering. Remarkable speedups are observed using the proposed GPU-based algorithm from experiments on routine medical data sets.

### 1 Introduction

Volume rendering deals with how a 3D volume is rendered and projected onto the view plane to form a 2D image. It has been broadly used in medical applications, such as for the planning of treatment [1] and diagnosis [2]. Unlike surface rendering, volume rendering bypasses the intermediate geometric representation and directly renders the volumetric data set based on scalar information such as density, and local gradient. This allows radiologists to visualize the fine details of medical data without prior processing such as the visualization of isosurfaces. Transfer functions are commonly employed for color mapping (including opacity mapping) to enhance the visual contrast between different materials. However, due to the large amount of computation involved, the rendering time increases dramatically as the size of the data set grows. Our objective is to provide radiologists with more efficient volume rendering tools to understanding the data produced by medical imaging modalities, such as Computed Tomography (CT) and Magnetic Resonance Imaging (MRI). Hence, we introduce several techniques, including hardware implementation using commercial graphics processing unit (GPU), to enhance the rendering speed. This increased speed will allow radiologists to interactively analyze volumetric medical data in real-time and in stereo.

According to [3], volume rendering approaches can be classified into three main categories: object-order, image-order and domain methods. Some hybrid

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methods [4] [5] are proposed by researchers in recent years, but their fundamental operations still fall into one of the three categories. The object-order approaches [6] [7] evaluate the final pixel values in a back-to-front or front-toback fashion, i.e., the scalar values in each voxel are accumulated along the view direction. Such intuitive approaches are simple and fast, but often yield image artifacts due to the discrete selection of projected image pixel(s) [8]. This problem can be solved by using splatting [9], which distributes the contribution of one voxel into a region of image pixels. While resampling in splatting is view-dependent, shear-warp [10] alleviates the complications of resampling for arbitrary perspective views. The input volume composed of image slices is transformed to a sheared object space, where the viewing rays are perpendicular to the slices. The sheared slices are then resampled and composited from front to back to form an intermediate image, which is then warped and resampled to get the final image. The image-order volume rendering [11] [12] is also in the category of ray casting or ray tracing. The basic idea is that rays are cast from each pixel on the final image into the volume and the pixel values are determined by compositing the scalar values encountered along the rays with some predefined ray function. One typical optimization is early ray termination, which stops tracing a ray when the accumulated opacity along that ray reaches a user-defined threshold. Another common optimization is empty space skipping, which accelerates the traversal of empty voxels. Volume rendering can also be performed in the frequency domain using Fourier projection-slice theorem [13]. After the volume is transformed from the spatial domain to the frequency domain, a specific 2D slice is selected and transformed back to the spatial domain to generate the final image. All of the previous three categories of methods can be partially or entirely implemented in GPU for acceleration [14] [15] [16]. Hardware-accelerated texture mapping moves computationally intensive operations from the CPU to the GPU, which dramatically increases the rendering speed.

A detailed comparison between the four most popular volume rendering techniques, *i.e.*, ray casting, splatting, shear-warp and 3D texture hardware-based methods, can be found in [17]. Experimental results demonstrate that ray casting and splatting generate the highest quality images at the cost of rendering speed, whereas shear-warp and 3D texture mapping hardware are able to maintain an interactive frame rate at the expense of image quality. When using splatting for volume rendering, it is difficult to determine parameters such as the type and radius of kernel, and the resolution of the footprint table to achieve an optimal appearance of the final image [8]. In shear-warp, the memory cost is high since three copies of the volume need to be maintained. The frequency domain methods perform fast rendering, but is limited to orthographic projections and X-ray type rendering [13].

## 2 Software-Based Accelerated Ray Casting

Software-based ray casting produces high-quality images, but due to the huge amount of calculation, the basic algorithm suffers from poor real-time performance. To accelerate software ray casting, there are two common acceleration techniques as mentioned in the previous section: empty space skipping and early ray termination.

Empty space skipping is achieved via the use of a precomputed min-max octree structure. It can only be performed efficiently when classification is done before interpolation, *i.e.*, when the scalar values in the volume are converted to colors before the volume is resampled. This often produces coarser results than applying interpolation first. If empty space skipping is applied with interpolation prior to classification, one additional table lookup is needed to determine whether there are non-empty voxels in the current region. Nevertheless, the major drawback with such kind of empty space skipping lies in that every time the transfer functions change, the data structure that encodes the empty regions or the lookup table needs to be updated.

Early ray termination exploits the fact that when a region becomes fully opaque or is of high opacity, the space behind it can not be seen. Therefore, ray tracing stops at the first sample point where the cumulative opacity is larger than a pre-defined threshold. The rendering speed is often far from satisfactory, even for medium-size data sets  $(e.g., 256^3)$ . In order to accelerate the speed, one can use acceleration techniques such as  $\beta$ -acceleration [18]. The fundamental idea of the  $\beta$ -acceleration is that as the pixel opacity (the  $\beta$ -distance) along a ray accumulates from front to back, less light travels back to the eye, therefore, fewer ray samples need to be taken without significant change to the final image quality. In other words, the sample interval along each ray becomes larger as the pixel opacity accumulates. Unlike  $\beta$ -acceleration, which depends on a pyramidal organization of volumetric data, here the jittered sample interval is applied directly to the data set. This reduces the computational cost of maintaining an extra data structure, especially when the transfer function changes. Instead of going up one level in the pyramid whenever the remaining pixel opacity is less than a user-defined threshold after a new sample is taken, the sample interval is modified according to a function of the accumulated pixel opacity:

$$s = s \times (1.0 + \alpha \times f) \tag{1}$$

where s denotes the length of the sample interval;  $\alpha$  denotes the accumulated opacity; and f is a predefined jittering factor. The initial value of s is set by the user. Normally, the smaller s the better image quality. For every sample point, the remaining opacity  $\gamma$  is compared against a user-specified threshold. If  $\gamma$  is less than the threshold, the current sample interval is adjusted according to Equation 1. We term this acceleration technique as  $\beta'$ -acceleration.

To further enhance the performance of software ray casting during interaction, the sample interval is automatically enlarged to maintain a high rendering speed, and once interaction stops, the sample interval is set back to normal. When multiple processors are available, the viewport is divided into several regions and each processor handles one region.

The whole process is executed in the CPU and main memory. The enhanced algorithm is illustrated in the following pseudo-code. Not only is this software

approach suitable for computers with low-end graphics cards, but since parallel ray tracing is used, it is also suitable for multi-processor computers or clusters.

Accelerated Software-Based Ray Casting

```
Break current viewport into N regions of equal size
Initialize early ray termination threshold \Gamma
Initialize jittering start threshold \Gamma'
Initialize jittering factor f
Initialize sample interval \boldsymbol{s}
For each region
  For every pixel in the current region
     Compute ray entry point, direction, maximum tracing distance {\cal D}
     While the traced distance d < D and \gamma < \varGamma
       Interpolate at current sample point
       Get opacity value \alpha according to opacity mapping function
       If \alpha \neq 0
          Compute pixel color according to color mapping function
          \gamma = \gamma \times (1.0 - \alpha)
       End If
       If \gamma < \Gamma'
          s = s \times (1.0 + \alpha \times f)
       End If
       d = d + s
       Compute next sample position
     End While
  End For
End For
```

## 3 GPU-Based Object-Order Volume Rendering

GPU-based object-order volume rendering has several advantages over GPU-based image-order volume rendering. First, perspective projections can be more easily implemented in object order, since only a proper scaling factor needs to be assigned to each slice based on several viewing parameters. In ray casting, the direction of each ray needs to be determined individually. Second, as pointed out in [19], GPU-based ray casting has the limitation that it can only render volumes that fit in texture memory. Since ray tracing needs to randomly access the whole volume, it is impossible to break the volume into sub-volumes and load each sub-volume only once per frame. Finally, most of the speedup from GPU-based ray casting comes from empty space skipping, and ray casting with only early ray termination shows close performance to object-order volume rendering while both implemented in the GPU, as compared in [15].

Other implementations generate the proxy polygons that textures are mapped to in the CPU, and use the fragment shader for trilinear interpolation and texture mapping. Little work has been done to exploit the vertex shader in the

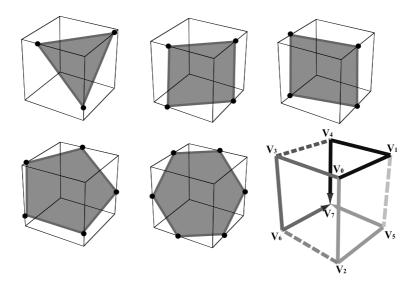


Fig. 1. The five intersection cases between a proxy plane and the volume bounding box, and the traversal order of the bounding box edges

hardware-accelerated volume rendering pipeline. Our accelerated rendering algorithm is based on the algorithm proposed by Rezk-Salama and Kolb [20], which balances the workload between the vertex shader and the fragment shader. Based on the observation of different box-plane intersection cases, the generation of proxy polygons can be moved from the CPU to the GPU. The intersection between a proxy plane and the bounding box of the volume may have five different cases, ranging from three intersection points to six, as illustrated in Figure 1. Let  $\mathbf{n} \cdot (x, y, z) = d$  represent a plane, where  $\mathbf{n}$  is the normalized plane normal and d is the signed distance between the origin and the plane, and let  $V_i + \lambda \mathbf{e}_{i,j}$  represent the edge  $E_{i,j}$  from vertex  $V_i$  to  $V_j$ , where  $\mathbf{e}_{ij} = V_j - V_i$ , then the intersection between the plane and the edge can be computed by

$$\lambda_{i,j} = \begin{cases} \frac{d - \mathbf{n} \cdot V_i}{\mathbf{n} \cdot \mathbf{e}_{i,j}}, \ \mathbf{n} \cdot \mathbf{e}_{i,j} \neq 0; \\ -1, & \text{otherwise.} \end{cases}$$
 (2)

If  $\lambda_{i,j} \in [0,1]$ , then it is a valid intersection; otherwise, there is no intersection. The edges of the volume bounding box are checked following a specific order, so that the intersection points can be obtained as a sequence that forms a valid polygon. If  $V_0$  is the front vertex (the one closest to the viewpoint) and  $V_7$  is the back vertex (the one farthest from the viewpoint), then the edges are divided into six groups, as shown in Figure 1 marked with different gray levels and line styles. For a given plane Pl parallel to the viewport that does intersect with the bounding box, there is exactly one intersection point for each of the three groups (solid lines), and at most one intersection point for each of the other

three groups (dotted lines). The six intersection points  $P_0$  to  $P_5$  are computed as described in Table 1. For the other seven pairs of front and back vertices, the only extra computation is to map each vertex to the corresponding vertex in this case, which can be implemented as a simple lookup table.

Table 1. The computation of the intersection points

Point	Checked	Edges	Intersection Position
$P_0$	$E_{0,1}, E_{1,4}$	and $E_{4,7}$	$\lambda_{i,j}$ , where $(i,j) \in \{(0,1), (1,4), (4,7)\} \land \lambda_{i,j} \in [0,1]$
$P_1$	$E_{1,5}$		$\begin{cases} \lambda_{1,5}, \lambda_{1,5} \in [0,1]; \\ P_0, & \text{otherwise.} \end{cases}$
$P_2$	$E_{0,2}, E_{2,5}$	and $E_{5,7}$	$\lambda_{i,j}$ , where $(i,j) \in \{(0,2), (2,5), (5,7)\} \land \lambda_{i,j} \in [0,1]$
$P_3$	$E_{2,6}$		$\begin{cases} \lambda_{2,6}, \lambda_{2,6} \in [0,1]; \\ P_2, & \text{otherwise.} \end{cases}$
$P_4$	$E_{0,3}, E_{3,6}$	and $E_{6,7}$	$\lambda_{i,j}$ , where $(i,j) \in \{(0,3), (3,6), (6,7)\} \land \lambda_{i,j} \in [0,1]$
$P_5$	$E_{3,4}$		$\begin{cases} \lambda_{3,4},  \lambda_{3,4} \in [0,1]; \\ P_4, & \text{otherwise.} \end{cases}$

In Rezk-Salama and Kolb's method, the coordinates of a sample point in the world coordinate system are required to be the same as the coordinates of the corresponding sample point in the texture coordinate system. However, this is not true for most cases, where the sizes of one volume are different in the two coordinate systems. The box-plane intersection test is carried out in the data coordinate system. Since typically the texture coordinates need to be normalized to the range between [0,1], a conversion of valid intersection points' coordinates is required. If the point  $P_k$  intersects the edge  $E_{i,j}$  at position  $\lambda_{i,j}$ , then each coordinate of the resulting texture-space intersection point  $P_k'$  is obtained by

$$P'_{k}.p = \begin{cases} \frac{V_{i.p-\min(B_{p})}}{\max(B_{p})}, \ e_{i,j}.p = 0; \\ \lambda_{i,j}, & e_{i,j}.p > 0; \\ 1 - \lambda_{i,j}, & e_{i,j}.p < 0. \end{cases}$$
(3)

where p denotes either x, y or z and B denotes the volume bounding box. The coordinates of  $P'_k$  are then scaled and translated in order to sample near the center of the cubic region formed by eight adjacent voxels in texture memory.

To further accelerate the rendering process, we also propose another enhancement that the sample interval is adjusted based on the size of the volume in the world coordinate system and the distance from the viewpoint to the volume. This idea of adaptive sample interval is similar to the concept of level-of-detail (LOD) in mesh simplification. The sample interval is calculated by:

$$s = S \times F^{\frac{\max(d)}{\max(B_x, B_y, B_z)}} \tag{4}$$

where S denotes the constant initial sample interval;  $F \ge 1$  denotes the predefined interval scale factor;  $B_x$ ,  $B_y$  and  $B_z$  denote the length of the volume bounding box B in the x, y, and z-direction respectively; and  $\max(d)$  denotes the distance between the farthest vertex of B and the view plane.

Now that the proxy polygons are generated, one can then perform texture mapping. The fragment shader performs two texture lookups per fragment to attach textures onto the proxy polygons. The first texture lookup gets the scalar value associated with the sample point from a 3D texture that holds the volumetric data. The hardware does the trilinear interpolation automatically for every sample. The second texture lookup uses the scalar value to get the corresponding color from a 2D texture that encodes the transfer function. Then, the textured polygons are written into the frame buffer from back to front to produce the final image.

The vertex program and the fragment program are both written in Cg, a high-level shading language developed by NVIDIA. To exploit the most powerful profile supported by a graphics card, the shader programs are compiled at runtime instead of at compile time. To accommodate graphics cards with different vertex processing capabilities, the amount of work assigned to the vertex shader should vary from card to card as graphic cards do not have all the same processing capabilities. The more capable the programmable graphics hardware is, the larger the amount of processing load is moved from the CPU to the vertex shader. Currently, our vertex program has variations for all the OpenGL vertex program profiles supported by the Cg compiler. The fragment program only requires basic Cg profiles to compile. Therefore, theoretically the proposed GPU-based volume rendering program can be executed on most commodity computers with a good-quality programmable graphics card.

#### 4 Results

The algorithms were tested on a dual-core 2.0GHz computer running Windows XP with a 256MB-memory NVIDIA GeForce 7800 GTX graphics card. The data used for testing is a medium-size (512x512x181) CT-scan of the pelvic region.

Software-based ray casting provides high quality images, but only with small viewports or for small data sets it can maintain an acceptable rendering speed, even with the proposed  $\beta'$ -acceleration. The rendering times using software ray casting with both early ray termination and  $\beta'$ -acceleration and with only early ray termination are enumerated in the first two columns of Table 2. Figure 2(a) depicts the two cases' performance curves with respect to the viewport size. The x-axis is the size of the viewport in pixels and the y-axis is the rendering time in seconds. The dark gray line denotes the performance of the method without  $\beta'$ -acceleration, and the other line denotes the performance of the one with  $\beta'$ -acceleration. On average, software ray casting with both early ray termination ( $\Gamma$ =0.02) and  $\beta'$ -acceleration ( $\Gamma'$ =0.6 and f=0.1) takes 28% less time than that with only early ray termination ( $\Gamma$ =0.02). The resulting images are shown in Figure 3(a)(b). There is no noticeable difference between these two images.

High-quality images and interactive rendering speed are both achieved by exploiting the processing power of the GPU. The rendering times under four

Viewport Size Rendering Time (unit: second) (unit: pixel) Without  $\beta'$ With  $\beta'$  Tex' 1.0 Tex' 1.15 Tex 1.0 Tex 1.15 200x200 0.172 0.125 0.031 0.016 0.015 0.015 300x3000.422 0.281 0.031 0.016 0.015 0.015 400x400 0.688 0.484 0.031 0.016 0.015 0.015 500x5000.750 0.047 0.031 0.015 1.078 0.031600x6001.641 1.125 0.0470.031 0.0310.016 1.562 700x7002.078 0.0620.0470.0470.031800x8002.704 2.187 0.0620.0470.0470.031 900x9003.391 2.469 0.0780.062 0.0620.0471000x10004.312 3.062 0.0940.078 0.078 0.047

Table 2. The rendering times using different acceleration techniques

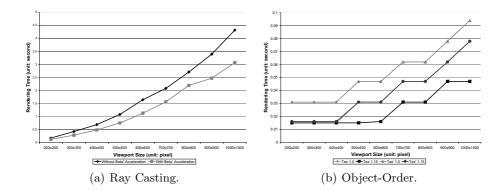


Fig. 2. The comparison of the rendering times using different acceleration techniques

different conditions are enumerated in Table 2. Tex' 1.0 denotes no acceleration; Tex' 1.15 denotes adaptive sample interval with interval scale factor F=1.15; Tex 1.0 denotes only with vertex shader acceleration; Tex 1.15 denotes with both acceleration techniques and F=1.15. Figure 2(b) gives a comparison of the performance curves under the four different conditions. In all cases, the rendering time increases as the viewport grows, but even for the 1000x1000 viewport the rendering times are below 0.1 second, i.e., the rendering speeds are above the psycho-physical limit of 10 Hz. With only adaptive sample interval enabled, when F=1.15, we get an average 33% speedup. With only vertex shader acceleration enabled, the algorithm's performance is almost the same as Tex' 1.15. With both acceleration techniques enabled, when F=1.15, an average 53% speedup is achieved with respect to the Tex' 1.0 case and an average 28% speedup is achieved with respect to the Tex' 1.15 case. The final images are shown in Figure 3(c)-(f), together with the images produced by software ray casting. From these images, no significant difference can be observed between the image quality of image-order methods and that of object-order methods, as long as the original data set is at high resolution.

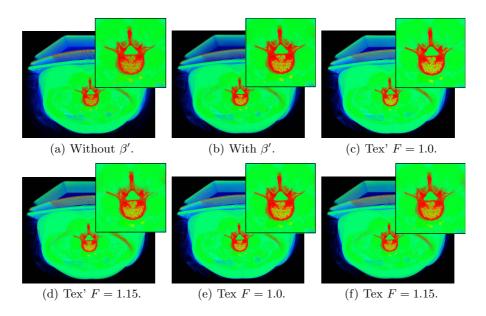


Fig. 3. Volume rendering results of a CT-scanned pelvic region

#### 5 Conclusion

In this paper, we have presented several volume rendering acceleration techniques for medical data visualization.  $\beta'$ -acceleration enhances the rendering speed of software-based ray casting using voxels' opacity information, while vertex shader proxy polygon generation and adaptive sample interval improve the performance of traditional hardware-accelerated object-order volume rendering. Remarkable speedups are observed from experiments on average-size medical data sets. We are now working on incorporating the  $\beta'$ -acceleration into the GPU ray casting pipeline, which may be more efficient than our current GPU-based object-order method. Moreover,we are also exploring more efficient and effective rendering algorithms using GPU clusters to handle larger and larger data sets produced by doppler MRI and temporal CT.

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