# **REMOTE RENDERING OF LARGE BIOLOGICAL DATASETS**

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Abstract: This paper presents remote rendering technologies to support the analysis and information exploration of large biological datasets. Reconstructed three-dimensional models and other data domains stored in a database are the foundation for the visualization. Remote Rendering via a clients web interface enables a widespread access to the data with interactive frame rates and high-quality rendering results without the demand of new client hardware. Moreover, the rendering server has the capability to manage multiple independent or collaborative sessions and has a very small bandwidth consumption over time.

## **1 INTRODUCTION**

Remote Rendering commonly means interactive visualization of three-dimensional datasets via a client/ server architecture over a network. Visualization of complex 3D models is still a task hard to accomplish without specialized hardware. Furthermore, ad-hoc visualizations over distance are playing a key role in future visualization trends. One problem of visualization is that huge amounts of data must be handled, which requires not only a fast CPU and high memory capacities (RAM), but also a high degree of graphics output generated by modern graphics cards.

These drawbacks can be overcome by the concept of remote rendering. The datasets are computed on a rendering machine with high graphics capabilities - the so-called *server*, and only rendered images are sent to the working station - called *clients*.

The work presented here is motivated by a given shortcomming regarding biological model visualization, namely limited bandwidth, slow clients with poor graphics capabilities, large biological 3D models and the demand to real-time interaction. The latter cannot be covered by existing conventional remote rendering techniques. Therefore, a remote rendering architecture with reduced network traffic generating correct high-quality renderings for low level end user PCs with interactive frame rates is given in this paper.

## 2 RELATED WORK

The idea of remote rendering is not new. Various remote rendering applications have been developed in the past. Adapted from (Schmalstieg and Gervautz, 1996), these techniques can be classified as follows:



Figure 1: Remote rendering types a) image-based, b) geometry replication, c) immediate-mode, and d) hybrid.

**Image-based.** Rendering is performed by the sender, and the resulting stream of pixels is sent over the network as provided in (Scheifler and Gettys, 1986) (see Figure 1 (a)).

Geometry-based. A copy of the geometric database

is stored locally to be accessed by the rendering process. The database can either be available before application start (kept on local hard disk), or downloaded just before usage, such as VRML / X3D browsers can do (Gueziec et al., 1999) (see Figure 1 (b)).

**Immediate-mode Drawing.** The low-level drawing commands used by drawing APIs are issued by the application performing the rendering. They are not immediately executed, but sent over the network as a kind of remote procedure call. The actual rendering is then performed by the remote machine (e.g. distributed GL (Shreiner et al., 2005)), see Figure 1 (c).

**Hybrid Rendering.** In this approach a 3D geometry as low resolution representation is rendered on the client's side and the server transmits images to the client as well. The used mode depends on the performed user action (computational vs. network load) (Koller et al., 2004) or is dynamically adapted to the available network capacities (see Figure 1 (d)).

Summarized, existing remote rendering techniques cover a wide range of problem statements including image-based remote renderings (suitable for fast network connections) or geometry-based remote renderings (e.g. cluster rendering with sufficient client's graphics hardware). Hybrid rendering methods for their capabilities could be identified as adequate rendering method for low and medium level clients using networks with medium network speed.

### **3 SYSTEM DEMANDS**

In this section the requirements of the involved underlying system architecture and its components as well as special user interests are discussed. Main issues are the individual clients' hardware conditions in the network, the data transfer capabilities and the trait of scientific data.

**Clients Condition.** A medium-sized scientific bio institution with approximately 300 potential clients was evaluated. The premises of these clients are completely heterogeneous, they cover a big inventory of out-of-date machines<sup>1</sup> as well as a few state-of-theart hardware machines. The present operating systems are as diverse as the clients' hardware condition. Whereas the processor speed is widely feasible, the graphics hardware is not. These preconditions have to be taken into account in the design of a remote visualization application for large biological models. **Server Condition.** Two different server machines could be used for testing the remote visualization performance.

The first one is a high performance computer with two Quad-Core Opteron 2.6 GHz, 16 GB RAM, a Quadro FX 5600 graphics card and furthermore a Tesla S870 processor with 500 GFlop and a shared 1 GBit connection (bottleneck) to the bio institution. An average server upload  $B_{up}$  of 2 MBit with average ping rates  $t_{avg}$  lesser than 23 ms could be identified  $(\sigma_{B_{up}} \approx 0.2 \text{ MBit}, \sigma_{tavg} 3 \text{ ms })^2$ .

The second server is a minimal graphics server configuration with a 2.8 GHz Pentium IV with 4 GB RAM. The rendering job is done by one *Nvidia 7800 GTX* graphics card. This server is directly located at the bio institution. The minimum network connection for this setup is bound to the local switches and achieves 100 MBit.

**Data Transfer.** The data transfer capabilities are one major aspect in the remote data visualization and can be formulated regarding to Equation 1.

$$B \ge \frac{(R_w R_h) C_{depth} f n}{c_{factor}} \tag{1}$$

Without any compression  $c_{factor}$  and for a screen resolution  $R_w x R_h$  (XGA), a color depth  $C_{depth}$  of 3 Bytes/pixel and a frame rate f of 26 frames/sec for only one client n, this would require at least a bandwidth B of:

$$B \ge \frac{(1024*768) \text{ pixel} * 3 \text{ Bytes}/\text{pixel} * 26 \text{ sec}^{-1} * 1}{1}$$

$$B \ge 60 MByte/s$$
 (2)

which is still high and only possible with at least 1 GBit/s ethernet<sup>3</sup> (see example calculation in Equation 2).

Using JPEG compression can reduce the image size by a factor  $c_{factor}$  of 10 - 300 (depending on the resulting image quality) and will also work with slower connections. The local network speed at the bio institution depends on the used switches and includes 100 MBit as well as 1 GBit connections. This fact prevents a standard remote rendering strategy due to the reason that high-quality renderings would not fit into a 100 MBit connection if a couple of other users share the same server connection (big *n*).

**Data Resources.** The biological data pool consists of different data domains from biological material. Available data include specifically microtome serial section data, gene expression assays utilizing mRNA

<sup>&</sup>lt;sup>1</sup>e.g. *Win98* Pentium III with less or equal 500 MB RAM

<sup>&</sup>lt;sup>2</sup>average measured with http://speedtest.com <sup>3</sup>SXGA requires around 100 MByte/s.

*in-situ hybridization* (ISH) probing of histological cross-sections as well as magnetic resonance spectroscopy measurements (<sup>1</sup>H-NMR) of caryopses at different points in time. Other potential datasets, such as macroarray gene expression data or metabolomic assays by laser microdissection (LMD), could be integrated into the database to enhance synergy and facilitate the inference of analysis results. Figure 2 presents a pie chart breaking down the data distribution per model.



Figure 2: Data distribution for one model, the absolute data totaling around 9 GB.

**Visualization Aspects.** Another very important precondition is the function volume of the remote rendering application to develop. At the moment the bio researchers test a software application which is especially tailored to them, namely the *BioVizIt* application. This software is initially a single-user stand alone application, which is not network compatible but covers the rest of the desired functionality.

All these features and actions have to be considered as part of the resulting remote rendering application to best possibly support the biologist's work.

As conclusion, there are different possibilities to perform remote visualization tasks as proposed in Section 2. As possible candidates only those solutions are appropriate which meet the requirements (recall Section 3).

As solely remote rendering approach which fits the system requirements a hybrid rendering approach is applicable. Due to the specific visualization requirements depicted in this section a conception with an explicit control to data exchange mechanisms and adaption possibilities of the visualization is necessary. A solution to achieve this is to extend the existing *BioVizIt* application with specific hybrid remote rendering capabilities.

# 4 THE HYBRID REMOTE RENDERING APPROACH

In the following, the remote visualization concept is presented, where a hybrid rendering approach is the basis. Furthermore, the architecture's adaptions of the BioVizIt platform, referring to network compatibility and approaches addressing limited bandwidth, are described.

**Hybrid Rendering.** The following Figure 3 illustrates the idea of the hybrid approach.



Image related Scene graph related Viewpoint related 3D related Session related

Figure 3: Sequential diagram of the proposed hybrid rendering approach.

In the beginning of the rendering process (initial rendering request) a request with datasets' names and initial settings is sent to the server. As a result, the server application is loading the requested file from the database, and in addition a low resolution 3D geometry model (*level of detail* also *LOD*) is transferred to the client. For optimization purposes the same geometry model is loaded only once. Moreover, the graphics server submits the first remote-rendered image to the client.

If in the following, user navigation is performed (consecutive rendering requests), the local 3D model is rendered by the client itself. Fast movements can be achieved without having performance problems due to network limitations. The local 3D model can be reduced up to one per mill of the original 3D model's data amount. Even if the bandwidth is less than 10 MBit an initial loading delay for an unoptimized data transfer of 4 seconds for the complete low resolution model would be acceptable to the user, if afterwards a stable real-time interaction can be performed. In case of stopping the navigation or changing the scene graph for example, a command is immediately sent to the server (containing at least the new camera position and orientation) to request a new rendered image. This approach ensures real-time model interactions with a high-quality rendering result during the model exploration.

*Geometry Submission:* The server stores the information of the original 3D models and also in coarser resolutions (*LOD*). Datasets are generated according to the proposed technique in (Schoor et al., 2009).

If a dataset is requested, the original dataset is loaded into the server application. At the same time a low resolution model is transferred from the database to the client.

*Image Submission:* The image submission by demand is a concept to overcome the server network load. Additionally, image compression algorithms *JPEG* 2000 (Taubman and Marcellin, 2001) are used to reduce the network traffic.

**Network Interconnect.** Another important aspect besides the underlying data transfer per se is how the data will be transported. To achieve high resolution rendering results on the client, only a reliable data transfer method is an option. This excludes for example *UDP* data transfer, because in this mode the correct rendering result is not warranted due to unordered or lost packages. The Transmission Control Protocol *TCP* is sufficient for the required purposes.

Adaption of the BioVizIt Platform. For an extension of the existing solution as server variant the functionality must be appended to network and session routines. Multiple network connections through clients must be promptly accepted and processed, using multi-threading techniques. User events are registered in a message queue. This queue is held in the operating system and members of the queue are passed to the application (for example panning the window). Received messages by the application are then handled in a procedure of the primary thread. To avoid blockages in the processing of this thread, a new thread - the "server thread" - is generated during the program start to wait for new client requests via TCP. If a client connects to the server over a predefined port, a new "client thread" is generated immediately for this client. This thread is responsible for further communication with the client.

The client thread reads the request from the server's socket thread and interprets it through the HyperText Transfer Protocol (HTTP). By accessing a virtual document, additionally the document area of the HTTP request is used, which includes the remote rendering control commands of the client for the server. These commands are processed sequentially.

At the end of this processing the client is enqueued to a list, which contains clients awaiting an image synthesis step.

### **5 RESULTS**

In this section, the essential aspects of the implementation are presented. Key issues for remote visualization are examined regarding the presented approach. **Latency of Network.** The latency at the beginning of a session depends on how the geometry transmission is handled. The coarse 3D model can first be accessed after full model upload. This can take a few seconds for the geometry transmission. This is an initial step which is only performed once a model is loaded. After that, the interaction is performed with latencies < 30 ms due to local rendering. The server rendering result takes less than 1 s after the mouse button is released (indicating the end of user navigation).

**Graphics Performance.** The graphics performance of the client rendering stage depends on its hardware and on the level of detail in which the 3D model was submitted. For the tested application even on a Pentium III client a coarse 3D model with 20,000 points could be interactively visualized. During the observation of the model no rendering step is required (no model interaction). That means that a real-time application can be guaranteed for the model interaction.

Network Throughput. The network throughput per high resolution image request (> 3-5 s) is equivalent to the physical size of this image and corresponds to approximately 300 kB. The geometry transfer per low resolution 3D model is approximately 4-5 MB in size. Scalability. The approach provided here has a good scalability. The rendering of scenarios, such as described in Section 3, can be accomplished on graphics servers without problems with interactive frame rates (Case 2: minimum graphics server solution). That means, in the worst case<sup>4</sup> the remote rendering image of the last handled client will be displayed with an additional delay of approximately 1 s if the data transfer is ignored. This is the theoretical maximum number of users which can be handled by the server under the assumption that a delay of more than 2 seconds is not acceptable to users. Taking the 100 MBit connection as lower bound for the data channel, more than 40 clients can request a full resolution image at the same time without causing additional delay at a client station. Assumed that a 2 MBit connection is the average connection condition for the server case 1, and a full image resolution is requested, only one client request per second is possible without causing an additional delay on the client.

**Error Recovery.** The error recovery is automatically performed by the network protocol layer (TCP) which provides explicit error corrections. Furthermore, a

<sup>&</sup>lt;sup>4</sup>e.g. a synchronous server request of 25 clients

transmission control and data ordering is also part of this protocol.

**Network Burst.** In general, the failure of a network connection in the beginning of a work session leads to an insufficient remote visualization. During the session a temporary burstiness of the network connection in the time scale of seconds can be bridged without problems due to the local downsampled version of the 3D model. A loss of connectivity within a larger time range disallows a practical use of the here provided technique due to the missing detail information the remote-rendered image provides.

**Security Issues.** The data pool of plant biological data consists of confidential information. Even an internal use must contain at least a simple protection mechanism due to project internal information. Therefore, a simple login query was integrated to authenticate the users with respective project rights. Furthermore, each session on the remote visualization server is initially hidden to other users.

**Balance between Local/Remote Resources.** The architecture of the hardware at the scientific institution is naturally very heterogeneous. A uniform strategy to extensively use clients' hardware is therefore not applicable. As the least common denominator, a low resolution 3D model, which is tailored to the actual hardware conditions, is used at the clients' side to achieve real-time interaction.

**Hardware/Software Incompatibility.** The visualization software is written in C++ and uses only platform-independent libraries. The visualization service tests for hardware premises, if specific features are not supported by the server hardware, alternatively CPU-based implementations exist (e.g. deformation model). The client's visualization front-end uses a platform-independent web interface, based on  $Java^{\text{(R)}}$  Version 1.5, Java  $3D^{\text{(M)}}$ .

Figure 4 shows a screenshot of the provided web interface and the visualization options.



Figure 4: Screenshot of the clients web interface showing a reconstructed model of a caryopsis with clipping plane.

#### 6 CONCLUSION AND OUTLOOK

In this paper a remote rendering approach was presented, which is applicable to medium-sized enterprises who have to deal with a large amount of scientific data. With a minimum of effort and the use of existing hardware this approach allows to visualize, explore and modify data. The clients hardware constitution must not be state-of-the-art which makes this approach of visualization furthermore attractive in a financial manner. The network bandwidth could be identified as limiting constraint respective to the number of clients contemporaneously used, therefore the local server with the minimal graphics configuration (case 2) is to prefer.

If the remote rendering application will be used with many users (> 30), the aid of more than one rendering machine can be useful. In the next term a larger focus will be placed on data modification possibilities. Another point is the popping effect which occurs by switching between the local LOD and the server image. This effect is not so perturbing as occurring in dynamic scenes. Anyway, a fading of the rendered image can reduce this effect.

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