

# High-Performance Computing for the Study of Earth and Environmental Science Materials Using Synchrotron X-ray Computed Microtomography

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Synchrotron x-ray computed microtomography (CMT) is a non-destructive method for examination of rock, soil, and other types of samples studied in the earth and environmental sciences. The high x-ray intensities of the synchrotron source make possible the acquisition of tomographic volumes at a high rate that requires the application of high-performance computing techniques for data reconstruction to produce the three-dimensional volumes, for their visualization, and for data analysis. These problems are exacerbated by the need to share information between collaborators at widely separated locations over both local and wide-area networks. A summary of the CMT technique and examples of applications are given here together with a discussion of the applications of high-performance computing methods to improve the experimental techniques and analysis of the data.

## 1. Introduction

Materials studied in the earth and environmental sciences are generally very inhomogeneous and complex materials. Investigation of the three-dimensional properties of these materials is essential. However, there are relatively few ways that these properties can be measured using non-destructive methods. These methods include laser confocal microscopy, magnetic resonance imaging, and x-ray computed tomography. The use of x-ray computed tomography is particularly powerful since it gives information on x-ray attenuation coefficients and thus can distinguish between different minerals, pore space, and liquid-filled pore space in specimens that can range from a few millimeters to many centimeters in size. Our purpose here is to describe the application of computed tomography techniques to these problems based on the use of synchrotron radiation and to discuss the application of high performance computing to improve the technique.

## 2. Instrumentation and Method

A schematic diagram of the CMT apparatus at the Brookhaven National Synchrotron Light Source (NSLS) is shown in Figure 1 [7]. The x rays are detected with a thin scintillator of CsI(Na) or YAG:Ce. A mirror/lens system is used to focus light from the scintillator onto a charge-coupled device (CCD) camera. Blurring effects caused by scattering of the x rays in the scintillator are minimized by the small depth-of-field of the magnifying lens. The spatial resolution of the system is energy dependent and

of the order of 0.005 mm. The CCD cameras used employed CCD chips with 1317 x 1035 and 3072 x 2048 pixels. In practice, the pixels are often binned to reduce the size of the tomographic volumes and thereby ensure practicable times for data acquisition and reconstruction. Monoenergetic beams are used at energies to about 20 KeV, and filtered white beam is used for measurements at higher energies to obtain higher x-ray intensities. Typical data collection times are of the order of 2-3 hours.

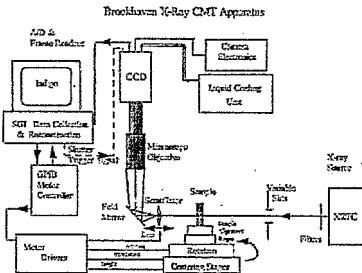


Figure 1. A schematic diagram of the apparatus.

Data acquisition produces a set of camera frames taken as the sample is rotated in a series of steps determined by the number of pixels desired in the volume. The procedure covers an angular range of 180 degrees with respect to the incident x-ray beam. The number and size of files depend on the sample size and desired spatial resolution.

The data reconstruction proceeds in 3 phases. Phase 1 applies a whitefield normalization, any filters needed, and writes files containing data from all views for a single slice (horizontal row of pixels), one file per slice. Phase 2 processes each slice independently. It applies the view-by-view air value normalization, optionally applies a filter to reduce the ring artifacts, computes the location in the images of the center of rotation, and converts the data to a sinogram. Phase 3 is the actual reconstruction. It generates a square array with dimensions of the horizontal row size. After this, the reconstruction is completed and the visualization process begins. This is a much more varied process and depends strongly on the particular sample being analyzed.

### 3. Experimental Results and Discussion

We have used the CMT apparatus to investigate many different materials relevant to the earth sciences. In particular, sandstones are of wide interest, and typical data is presented here to show a specific example of the usefulness of synchrotron CMT.

The data can be presented in several ways. A volume representation showing a 3-D view of sandstone from the Vosges region of France is shown in Figure 2. The grain structure of the material is clearly visible. A view of a single section through the sample is shown in Figure 3. This type of view helps to highlight the pore-grain relationships.

The data can then be processed according to the measured attenuation coefficients to display the data in binary form representing either solid or pore space. Analysis of this data gives the two-dimensional correlation function, porosity, permeability, and tortuosity. The measured microgeometry also can be used as a realistic basis for fluid flow calculations [2].

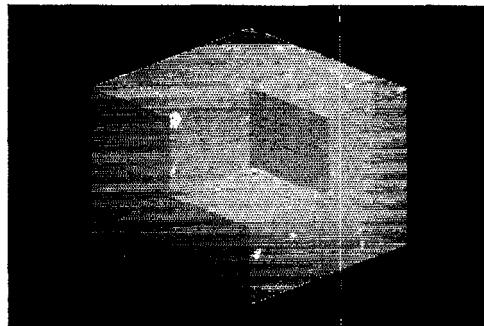


Figure 2. Isosurface rendering of a three-dimensional tomographic volume of a sample of sandstone from the Vosges. The color scale indicates the values of the measured absorption coefficients. The pore space is shown as blue.

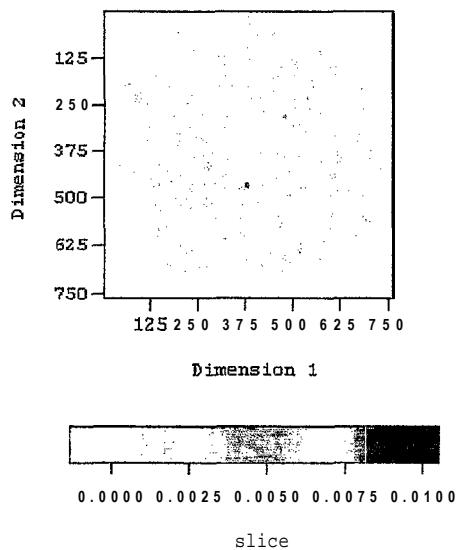


Figure 3. Single section through the sandstone volume shown in Figure 2. Absorption coefficients are indicated by the color scale.

#### 4. High Performance Computing

The present system used at BNL and others, similar in design at the NSLS, APS, ESRF, and other synchrotrons, have demonstrated their worth in varied experiments on environmental and earth science topics. However, the system performance and usefulness can be vastly improved by use of high-performance computing technology.

First, present day CCD cameras can produce 15-20 frames per second with a size of about 1000 x 1000 pixels. Assuming that about 1500 frames will be necessary to acquire data for a tomogram, approximately 6 gigabytes of data are produced per second (2 bytes per pixel). The goal is then to carry out all steps of the reconstruction process in the time needed to acquire the data, or about 1500 frames/15 frames/s or 100 s.

Second, in order to be able to adjust experimental parameters in near real time, it is necessary to produce data displays on about the same time scale. There needs to be control of the display by the experimenter so that **different** views are feasible and **thresholding** can be set to display changes in pore structures or fluid motion.

There are other aspects of the problem that **include** high-speed networking, rapid data storage, and remote viewing that present challenges to computer science. Here we concentrate our attention on the **first** two points, data reconstruction and viewing. We have developed methods for doing these tasks on the time scales required by recourse to parallel computation techniques.

X-ray computed microtomography is a highly computer intensive and memory intensive **application**. The large volumes and small grid spacing required for micrometer resolution push the limits of even the most **powerful** workstations. For this reason we are applying high **performance** parallel computation and data compression for remote access of tomographic data sets. The two main areas that need to be addressed are reconstruction and visualization. Reconstruction takes many projections obtained **from** the high resolution **CCD** camera and uses a FFT transform method developed by Robert **Marr** at **Brookhaven** to form a three-dimensional gridded data set. Visualization **of the** data set either as a volume or an **isosurface** is used to closely inspect the **sampb** and extract new features. This is typically done within a complete visualization package such as **OpenDX** or **VTK**.

Reconstruction of large data sets can take several hours. **Visualization** on large grids can also take tens of minutes just to form a particular view of an **isosurface**. The use of **parallelization** to address these problems is one way to achieve necessary speedups for **fast** reconstruction and **visualization**. Basically one divides up the data set across multiple processors and reassembles the **final** result by synchronization and communication among processors. There are two main protocols for **parallelization**: **Parallel Virtual Machine (PVM)** and the **Message Passing Interface (MPI)**. **MPI** is the more recent of the two protocols and is becoming a standard particularly on clusters of Linux computers. Parallel reconstruction **using MPI** is currently being used at the **APS**, and this technique **can also** be applied to data **from** the **NSLS** beam line. The results of applying parallel **visualization** using **MPI** to the **NSLS** beam line data are **described** here.

## 4.1 Parallel Visualization

**Parallel** visualization refers to the use of multiple computers for the graphical depiction of large data sets. It has its origins in parallel rendering where one breaks up the **image** to **be** rendered into smaller pieces and has each processor render its own part with **all** the pieces brought together for the final composite image [6, 8]. For example, Baily [5] has **ported** Pixar's **RenderMan** rendering **software** to the parallel Intel paragon machine. The **free** ray tracing program **POV** has also been ported to Linux clusters using **MPI** and **PVM** and achieved near linear speedups comparable with more expensive specialty supercomputers. Recently, parallel visualization has gone a step beyond this by **breaking** up the data set itself. This data parallel model allows **large** data sets of high **resolution**, such as the x-ray tomography data sets discussed here, to be manipulated by using the cumulative memory and processing power of a Linux cluster. The data parallel model has been **implemented** in **OpenDX** 4.1.2 and **VTK** using **MPI** [1, 3].

### 4.1.1 OpenDX

**OpenDX** is an **open source visualization package** that can be applied to a wide variety of data sets [1]. Currently, we use the software to give a quick view of x-ray tomographic data sets, **extract**

isosurfaces and slices, and convert from NetCDF data format to a data format that can be read into VTK. Recently, a port of the software using MPI has been achieved, and we are planning to apply this version to x-ray tomography. Figure 5 shows a screen save of the OpenDX software we have developed. OpenDX builds applications by drag& modules of specific functionality into the visual programming editor canvas and linking them together to form a network. The network shown uses an isosurface module, an export module to do the NetCDF conversion, and an image module for the final rendering. The data set shown is a high resolution x-ray tomographic data of the thigh bone of a rat used in osteoporosis studies.

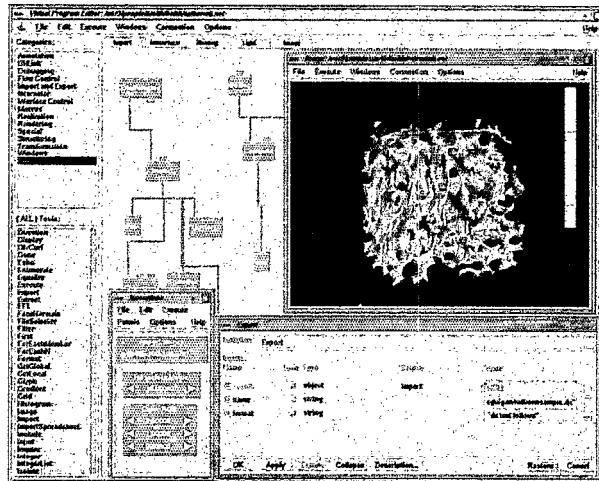


Figure 5. OpenDX application for isosurface extraction and conversion from NetCDF data format to VTK format.

### 4.1.2 VTK

The ‘Visualization Toolkit (VTK)’ is a **multiplatform** visualization package with C++, TK/TCL and Python **bindings**. It is known for its high quality implementation of the latest algorithms in computer graphics. Its **most** recent version includes a **parallel MPI implementation** of a subset of its **modules** [3, 4]. Here we discuss three programs built on this **subset** which we have applied to x-ray tomographic data S&S.

The first program, **Paralso**, breaks up the data set according to the command line arguments. It then computes isosurfaces for each piece on separate processors and then renders the final image with colors to indicate the work of the separate processors. An example for a tomographic data set is shown in Figure 6. The computation was done on a 4 Xeon processor 1400 Linux server from SGI.

The second MPI program **DataParallelism** breaks up the data more directly by using an image reader module. It includes its own **sample data and explicit timing functions** to measure the performance of the parallel isosurface computation. The results for the sample data are shown in Figure 7 and indicate nearly a factor of three speedup using all four processors.

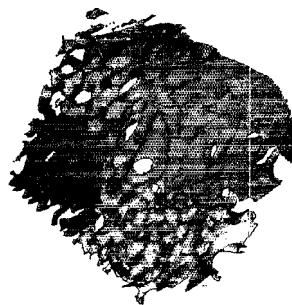


Figure 4. Output from the VTK program **ParaIso**. The four colors indicate the portion of the **isosurface** computed by each of the four processors on the **Linux** cluster.

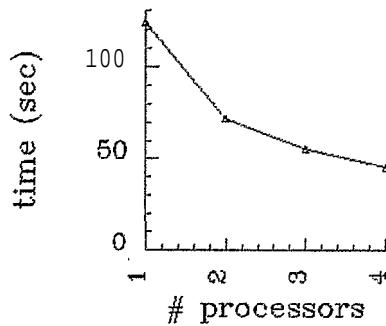


Figure 7. Graph showing the speedup using the parallel MPI VTK software **DataParallelism** to compute **isosurfaces**.

The third program, **ParaView**, is potentially the most powerful of the three. It includes a **Tk** widget control panel for **interacting** with the parallel MPI VTK application. This includes functionality to **input** various data sources, control over **isosurface** values, and **thresholding** for volumetric data. We are currently **modifying** the code to directly input x-ray tomographic data. A screen shot of the **user interface** applied to sample data is shown in Figure 8.

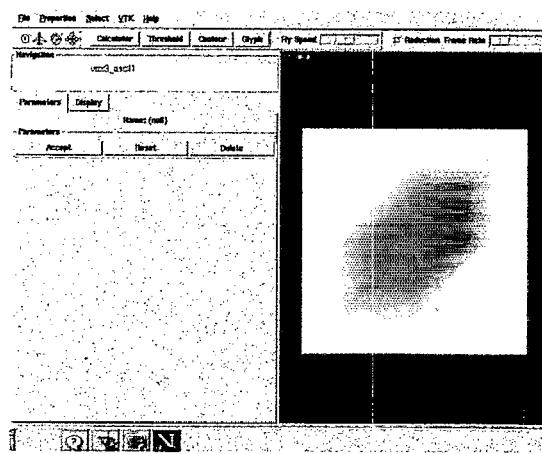


Figure 8. User interface for the parallel MPI VTK program, **ParaView**, shown with sample data set.

## 4.2 Remote Visualization

The use of large computer **facilities** to perform x-ray tomographic reconstruction and visualization is of **no** use unless the final rendering can be delivered to the scientist's desktop. This can mean delivery over a high-speed intranet or more remote delivery connecting dispersed researchers over the Internet or wide area network. We are currently testing two software packages designed to perform remote visualization. Both packages use compression algorithms to speed up transmission to the desktop.

The **first** remote visualization software, **VizServer** from **SGI**, transmits **OpenGL** visualization remotely **from** a multipipe **SGI Onyx2** computer to **SGI**, **Sun**, or **Linux** clients. The **software** was able to achieve frame rates **useful** for interactive viewing (approx. 10 **frames/sec**) for typical data sets. The **Onyx2** we used had 2 **GB** of Ram., 6 processors and 2 pipes. As only one pipe is available for **vizserving**, this means at most one user at a time can access the data remotely **from** this system

A second remote visualization software has recently been released by **TGS** as part of **OpenInventor** 3.0. **OpenInventor** is a high level **C++** and **Java** graphics toolkit built on top of **OpenGL**. This software works on a variety of **platforms** and doesn't require separate pipes for each user. It delivers **OpenInventor** applications directly to the desktop, and these can include x-ray tomographic data in **Inventor** and **VRML** format.

Another approach to remote visualization is the construction of a **VRML** server to deliver 3-D models directly over the Internet. **VTK** itself can be used to construct such a **server** in conjunction with a **VRML** browser. Currently we are using **Cosmoplayer** and **parallelgraphics** **VRML** browsers. The later can deliver on a variety of platforms **including** wireless **PDA** devices.

## 4.3 Stereoscopic Viewing

To understand the three-dimensional structure **from** high-resolution x-ray computed tomography it is **useful** to have a stereoscopic presentation of the data. Currently we are **using** a passive system with two **Barco** projectors and polarized **filters** [9]. The projectors are connected to an **Onyx2** computer and the visualization is rear projected on a special screen that preserves polarization. The final visualization is suitable for group viewing in a visualization theatre with inexpensive polarized glasses. At the desktop, stereoscopic viewing can be done using a page-flipping method and active **glasses**. Both methods can be used with **OpenDX** or **VTK** visualization toolkits. The parallel graphics **VRML** browser can also be operated for remote stereoscopic visualization on the Internet.

## 5. Conclusions

In this paper we briefly **described** the use of **synchrotron CMT** for investigation of earth and environmental sciences samples and then described improvements to the **CMT** system by application of high performance computing methods. We used **OpenDX** and **MPI** versions of **VTK** to perform the visualizations and achieved a **factor** of three increase in speed using a four processor **Linux** cluster. We also studied the application of visualization server software for desktop delivery and achieved **reasonable** **frame** rates for typical data sets. In the **future**, we will apply **MPI** versions of **OpenDX** and the use of **VRML** servers in order to make these high performance techniques widely available.

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