GPU-Accelerated SART Reconstruction Using the CUDA Programming Environment

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ABSTRACT

The Common Unified Device Architecture (CUDA) introduced in 2007 by NVIDIA is a recent programming model making use of the unified shader design of the most recent graphics processing units (GPUs). The programming interface allows algorithm implementation using standard C language along with a few extensions without any knowledge about graphics programming using OpenGL, DirectX, and shading languages.

We apply this novel technology to the Simultaneous Algebraic Reconstruction Technique (SART), which is an advanced iterative image reconstruction method in cone-beam CT. So far, the computational complexity of this algorithm has prohibited its use in most medical applications. However, since today’s GPUs provide a high level of parallelism and are highly cost-efficient processors, they are predestined for performing the iterative reconstruction according to medical requirements.

In this paper we present an efficient implementation of the most time-consuming parts of the iterative reconstruction algorithm: forward- and back-projection. We also explain the required strategy to parallelize the algorithm for the CUDA 1.1 and CUDA 2.0 architecture. Furthermore, our implementation introduces an acceleration technique for the reconstruction compared to a standard SART implementation on the GPU using CUDA. Thus, we present an implementation that can be used in a time-critical clinical environment.

Finally, we compare our results to the current applications on multi-core workstations, with respect to both reconstruction speed and (dis-)advantages. Our implementation exhibits a speed-up of more than 64 compared to a state-of-the-art CPU using hardware-accelerated texture interpolation.

Keywords: CT, CTCB, RECON, CUDA, GPU, SART, OTHER

1. INTRODUCTION

SART\textsuperscript{1} is a well-studied reconstruction method for cone-beam CT scanners. Already ten years ago it was obvious that, in certain scenarios, the Algebraic Reconstruction Techniques (ART) has many advantages over the more popular filtered back-projection\textsuperscript{2} (FBP) approaches. Due to its higher complexity, ART is rarely applied in most of today’s medical CT systems. The typical medical environment requires fast reconstructions in order to save valuable time. Industrial solutions face the performance challenge using dedicated special-purpose reconstruction platforms with digital signal processors (DSPs) and field programmable gate arrays (FPGAs). The most apparent downside of such solutions is the loss of flexibility and their time-consuming implementation, which can lead to long innovation cycles. In contrast, the Common Unified Device Architecture (CUDA) enables a conspicuous simplified development with improved but still limited flexibility. We have already shown that current GPUs offer massively parallel processing capability that can handle the computational complexity of three-dimensional cone-beam reconstruction.\textsuperscript{3} In 2007 NVIDIA introduced the QuadroFX 5600 architecture (359.3 GFlops\textsuperscript{*}, 1.4 GHz, one multiply-add operation per clock cycle per Stream Processor), which uses 128 Stream Processors in parallel.

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\textsuperscript{*}1 GFlops = 1 Giga floating point operations per second
Since then, NVIDIA has developed and introduced new CUDA compatible computing architectures with even higher peak performance for solving complex computational problems on the GPU. Recently NVIDIA introduced the QuadroFX 5800 architecture offering 240 Stream Processors for parallel computation. In August 2008 NVIDIA released the second major release of their programming model, CUDA 2.0, enhancing flexibility and extending the hardware feature support.

So far, the iterative reconstruction performance on graphics accelerators has often been evaluated using graphics-based implementation approaches involving OpenGL and shading languages. CUDA offers a unified hardware and software solution for parallel computing on CUDA-enabled NVIDIA GPUs supporting the standard C programming language together with high performance computing numerical libraries†. In this paper we describe our approach for dividing the computational steps and we measure the reconstruction performance using this innovative programming model.

Originating from ART, SART is an iterative method to reconstruct a volumetric object from a sequence of alternating volume projections and corrective back-projections. By measuring the difference between the current volume’s forward-projection and the projections acquired by the scanner, a corrective image can be distributed onto the volume grid in the back-projection step. Repeating this correction procedure makes the volume fit to almost all projections or at least minimizes the error in the case of convergence.

Differing from the original ART, where the volume is corrected ray by ray, SART performs a projection-wise correction. This concept becomes the SIRT method when taken one step further to compute all the correction images before they are back-projected onto the volume grid. Ordered Subsets (OS) approaches are in between SART and SIRT, because they compute a set of corrective images and back-project them before computing the next set of corrective images. Therefore, the projections are partitioned into disjoint sets.

ART-based methods have many advantages over the commonly used FBP and FDK methods for 3D cone-beam reconstruction when a full set of projections is not available (e.g., short scan CT or super-short scan CT) or the projections are not distributed uniformly in angle. Furthermore, when projections are sparse or missing at certain orientations, iterative methods can provide superior image quality. If an object is extended beyond the scanning field of view, satisfactory reconstruction results can be produced from incomplete data by applying iterative reconstruction techniques with special constraints.

Comparing our results to other approaches towards accelerating iterative CT-image reconstruction, Xu et al. recently published a similar approach. In their paper the authors report on the efficiency of GPU accelerated iterative reconstruction. Therefore they insist of a general API to use GPUs, neglecting if the API is used for CG or CUDA implementation. Due to the fact of their statement that a ‘write operator can be executed only at the end of a computation’, it is clear that they used OpenGL and shading languages as before. With CUDA as the API of choice, this isn’t the case. CUDA allows write operations to the global memory during a kernel execution which is used in our back-projection implementation. In Xu’s approach the volume is represented as a stack of slices (2D arrays) analogous to our CUDA 1.1 approach. Despite from OpenGL and CUDA they have already shown an improved time performance on GPUs for an OS approach compared to the SART method.

In this work, we evaluate the performance of a more advanced reconstruction approach on the current GPU technology. We consider the parallelized implementation and optimization of the SART method for CUDA 1.1 and CUDA 2.0 and introduce a new concept for improving iterative reconstruction speed on GPUs. In the following Section we will briefly review the SART reconstruction theory and describe our CUDA implementation approaches in detail. In Section 3 we will report about the exemplary experiment and the achieved results. Finally, we will conclude our work.

2. METHODS AND IMPLEMENTATION

Algebraic Reconstruction Techniques in principle solve a system of linear equations according to the Kaczmarz method. As introduced SART performs a projection-wise correction of the object.

The SART method can be divided into two steps: the corrective image computation (including the forward-projection step, difference computation and scaling) and the back-projection step. In the latter part we use

†http://www.nvidia.com/cuda
Figure 1. Perspective geometry of the C-arm device (the z-axis and z-axis are not necessarily parallel) together with
the parallelization strategy of our back-projection implementation on the GPU using CUDA (the x-z plane is divided
in several blocks to specify a grid configuration, and each thread of a corresponding block processes all voxels in y-direction).

our voxel-driven method\(^3\) in order to distribute the corrective image onto the volume. For the corrective image
computation a forward-projection based on ray casting\(^{11}\) is employed to compute projections of the current
volume estimation. The computed projections are then compared to the acquired projections. We adapted
our implementation of a CUDA-based ray casting algorithm,\(^{12}\) where we make use of the hardware-accelerated
interpolation of textures on graphics cards to compute the trilinearly interpolated samples along each ray.

In theory, the system matrix for the SART defines the forward- and back-projection such that the methods
are transposed to each other. In our case, we have an unmatched forward-/back-projector pair for the iterative
reconstruction, which has already been investigated.\(^{13}\) The authors demonstrate that an unmatched pair, where
the forward-projector is ray-driven while the back-projector is voxel-driven, can effectively remove ring artifacts
compared to a matched pair.

Because of deviations due to mechanical inaccuracies of real cone-beam CT systems such as C-arm scanners,
the mapping between volume and projection image positions is described by a \(3 \times 4\) projection matrix \(P_i\) for
each X-ray source position \(i\) along the trajectory. In a calibration step the projection matrices are calculated
only once when the C-arm device is installed. As we use a voxel-based back-projection approach it is required to
calculate a matrix vector product for each voxel and projection. However, for neighboring voxels it is sufficient
to increment the homogeneous coordinates with the appropriate column of \(P_i\).\(^{14}\) The homogeneous division
is still required for each voxel. It cannot be neglected for a line parallel to the z-axis of the voxel coordinate system
since, in general, the projection planes are slightly tilted with respect to this axis (see Figure 1 for an illustration
of the perspective geometry of a C-arm device). Due to the fact that the ray direction can be extracted out
of the projection matrix, the calibrated geometry is also used in the forward-projection step. We intentionally
avoided to use detector re-binning techniques that align the virtual detector to one of the volume axis since this
technique decreases the achieved image quality and requires additional computations for the re-binning step.\(^{15}\)

One of the important changes between CUDA 1.1 and CUDA 2.0 is the support of 3D textures. We will first
review the implementation of the back-projection step, because this is not affected by different CUDA versions.
After that we detail the forward-projection step which can exploit the new features of CUDA 2.0.

**Back-projection**

The back-projection step performs a voxel-based back-projection, which requires the calculation of a matrix-
vector product for each voxel in order to determine the corresponding corrective projection value. We invoke our
back-projection kernel on the graphics device, where each thread of the kernel computes the back-projection of
a certain column and volume slice (see Figure 1).
This allows to save six multiply-add operations by incrementing the homogeneous coordinates with the appropriate column of \( P_i \) for neighboring voxels in \( y \)-direction. This approach also reduces the register usage (see Figure 2). In this regard, a rectification-based approach has theoretically the potential to further improve the reconstruction speed.\(^{15}\) During our experiments we, however, observed that this is not true for the used graphics hardware. The rectification-based method approach even degrades the reconstruction speed when compared to the approach described above.

An efficient grid configuration for the back-projection step was chosen experimentally for an arbitrary back-projection configuration (see Figure 3). A more detailed description can be found in Scherl et al.\(^3\)

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**Figure 2.** Dependency of the multiprocessor warp occupancy on the register usage. Our CUDA back-projection implementation uses only 10 registers.

**Figure 3.** Back-projection execution time example for different grid configurations.

Corrective image computation

As introduced SART performs a projection-wise correction of the current estimation of the volume. Therefore, the corrective image has to be computed from the difference between the original projection and the appropriate simulated X-ray image of the current reconstruction estimate. All values in the corrective image are finally multiplied by the relaxation factor before the back-projection step. The implementation principles for CUDA 1.1 and CUDA 2.0 are illustrated in Figures 6 and 7.

A realistic simulation of the X-ray imaging process can be achieved by a ray cast based forward-projection. Research on this grid-interpolated scheme, where the interpolation is performed using a trilinear filter and the integration according to the trapezoidal rule, showed that the root mean square (RMS) error is comparable to other popular interpolation and integration methods used in computed tomography. This scheme is our first choice, because it can be ideally mapped to the GPU hardware including hardware-accelerated texture access.

**Algorithm 1** Forward-projection with a ray casting algorithm

```plaintext
for all projections do
    compute source position out of projection matrix
    compute inverted projection matrix
    for all rays inside the projection do
        compute ray direction depending on the image plane
        normalize direction vector
        //RAY CASTING
        compute entrance and exit point of the ray to the volume
        if ray hits the volume then
            set sample point to the entrance point
            initialize the pixel value
            while sample point is inside the volume do
                add up the computed sample value at current position to the pixel value
                compute new sample point for given step size
            end while
        else
            set pixel value to zero
        end if
    end for
end for
```

The volumetric ray casting principle for the forward-projection step is illustrated in Figure 4 and the algorithm is shown in Algorithm 1. To determine the attenuation value of a certain pixel on the detector plane, a ray is drawn pointing from the X-ray source towards the detector pixel position. Afterwards voxel intensity values inside the volume are sampled equidistantly along the ray. These sampling values add up to the respective
attenuation value in the simulated projection. Similar to the back-projection step we use projection matrices, instead of assuming an ideal geometry, to compute the resulting perspective projection.

To parallelize the forward-projection step, each thread of the kernel computes one corrective pixel of a projection. Analogous to the back-projection step we chose the grid configuration experimental due to our results. In the implemented kernel we compute the direction vector for a specific ray, which is the first step in the inner for loop in Algorithm 1. Therefore we take the source position vector and the 3D coordinate of the pixel position, compute the difference vector, and normalize it. The source position for all rays of a projection is obtained from the homogeneous projection matrix which is designed to project a 3D point to the image plane. Depending on the output format of the projection (2D image- vs. 3D world-coordinates), this matrix has three or four rows. In the latter case, the vector can be found in the fourth column of the inverted matrix (first three components). In the case of a $3 \times 4$ matrix it is possible to drop the fourth column, invert the $3 \times 3$ matrix and multiply the inverse with the previously dropped fourth column to get the source position. This holds, because in case of a perspective projection with projection matrices, this fourth column represents the shift of the optical center to the origin of the coordinate system. Galigekere et al. have shown already how to reproject using projection matrices.

![Figure 5. GPU implementation principle: Volume represented in a 2D texture by slices $S_j$ is forward-projected (FP). After computing the corrective image and scaling with the relaxation factor, the back-projection (BP) distributes the result onto the volume. After performing an update the 2D texture representation of the volume is equal to the volume.](image)

In the kernel code, the inverse of the projection matrix is used to get the ray direction out of the pixel position in the projection image. The entrance and exit positions of the specific ray into the volume are calculated and stored as entrance and exit distances with respect to the source position. Between those points the volume is then sampled equidistantly. To get one sampling position, we take the entry vector and add the direction vector multiplied with the step size times a counter variable. The following sampling step itself proves to be crucial for the algorithm’s efficiency. In order to get satisfying results, a sub-voxel sampling is required, which introduces a trilinear interpolation.

The global memory offers write access and thus has a higher latency. In contrast read-only texture memory has conspicuous low latency due to caching mechanisms and further offers hardware-accelerated interpolation. In CUDA 1.1 the computation of each sample point intensity is a critical issue since support for 3D textures is not provided. In consequence, a workaround had to be applied that used just the bilinear interpolation capability of the GPU. The kernel computes a linear interpolation between stacked 2D texture slices ($S_j$) (see Figure 6). Therefore, two values are fetched from proximate stack slices with hardware-accelerated bilinear interpolation and afterwards linearly interpolated in software. These sampling steps are substituted by only one hardware-accelerated 3D texture fetch in CUDA 2.0. Since texture memory is read-only, the back-projection updates the
original volume data kept in global memory. The volume-representing texture has to be synchronized with the updated estimate (Figure 6). Such a synchronization is referred to as a texture update.

![Figure 6. GPU implementation principle: Volume represented in a 3D texture is forward-projected (FP). After computing the corrective image and scaling with the relaxation factor, the back-projection (BP) distributes the result onto the volume. After performing an update the 3D texture representation of the volume is equal to the volume.](image)

The difference in volume representation for the corrective image computation leads to two major principles of SART implementation using CUDA shown in Figure 6 for CUDA 1.1 and Figure 7 using CUDA 2.0. After all corrective images have been computed and back-projected for all iterations the reconstruction finishes by transferring the volume to the host system memory.

### 3. EXPERIMENTS AND RESULTS

In order to evaluate the performance of the GPU vs. the CPU we did the following experiment. On the CPU side we used an existing multi-core based reconstruction framework, while using NVIDIA’s QuadroFX 5600 on the GPU side. Our test data consists of simulated phantom projections, generated with DRASIM. We used 228 projections representing a short-scan from a C-arm CT system to perform iterative reconstruction with a projection size of 256 × 128 pixels. The reconstruction yields a 512 × 512 × 350 volume. In order to achieve a sub-voxel sampling in the forward-projection step we used a step-size of 0.3 of the uniform voxel-size. Since the majority of time in reconstruction is spent on copying the volume data for the reconstructed image from the global memory to a texture memory in order to use the hardware-accelerated interpolation, we can significantly reduce this time by performing an ordered subsets method.

Table 1 shows the achieved performance for the CPU-based SART reconstruction as well as for our optimized GPU implementations using CUDA 1.1 and CUDA 2.0. The former does not need additional memory for the forward-projection step because there is no texture update, and therefore reconstruction times for the SART and OS are identical.

In principle, graphics cards have a very high internal memory transfer rate (≈ 62 GB/s on the QuadroFX 5600). Since texture memory is not stored linearly, it has to be reorganized for texture representation, which is the rate-limiting factor using CUDA 1.1. We measured 476 seconds to transfer a 512³ volume 414 times to the texture stack representation. This is approximately 1.15 seconds for a single texture update. Using 3D textures in CUDA 2.0 this can be improved by a factor of 10 such that a texture update can be performed in approximately 0.11 seconds.
A SART implementation in CUDA 1.1 wastes approximately 1 second per update to transfer the back-projected volume to a texture memory representation used for the forward-projection. If the number of forward- and back-projections between two texture updates is increased slightly, the reconstruction speed will be faster while the convergence rate remains almost at the same level, but is not as fast as a standard SART (analogous to the relation between ART and SART). This trade-off between convergence and speedup is one of our main results. The convergence was recently examined by Xu et al.\textsuperscript{10}

We compared reconstruction times on three different systems. First, an off-the-shelf PC equipped with an Intel Core2Duo processor running at 2 GHz, second, a workstation with two Intel Xeon QuadCore processors at 2.33 GHz and a NVIDIA QuadroFX 5600 with CUDA 1.1 and 2.0. The SART implementation using CUDA 1.1 is the slowest implementation on the GPU. Yet it is more than 7.5 times faster than the PC and 50 percent faster than the workstation. Employing the ordered subsets optimization yields another speedup of over 4 times. SART with 3D texture interpolation (CUDA 2.0) is even a bit faster, and using OS again results in a total speedup of 64 and 12 compared to the PC (2 cores) and the workstation (8 cores) respectively.

### 4. CONCLUSION

In conclusion, we have optimized the reconstruction speed and the convergence behavior in our algorithm design. We have shown the advantage of using the texture memory of current graphics cards to perform the most time-consuming parts of an iterative reconstruction technique effectively on the GPU using CUDA 1.1 and recently released CUDA 2.0. Apparently, in CUDA 1.1 the time consuming texture updates dominate the overall reconstruction time. This is dramatically relieved in the CUDA 2.0 implementation. Therefore, the impact of OS with CUDA 2.0 is much lower than with CUDA 1.1.

Furthermore, we have demonstrated the drawback of representing the volume data as a texture in that, it is the time-expensive update process, necessary in iterative reconstruction. Alternatively, the OS method reduces this effect because it requires fewer updates. For a small increase of forward-/back-projections between two update steps, the reconstruction speed is accelerated significantly, while the convergence rate is not decreased significantly. Due to a reconstruction time of less than 9 minutes, our implementation is already applicable for specific usage in the clinical environment.

### 5. OUTLOOK

Our research also demonstrates that there exists a lack of comparibility for fast 3D reconstruction implementations, despite the plurality of publications. For future research, we want to improve this by providing an open platform –RabbitCT (www.rabbitCT.com)– for worldwide comparison in back-projection performance on different architectures using a specific high resolution angiographic dataset of a rabbit. This includes a sophisticated interface for rankings, a prototype implementation in C++ and image quality measures.
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REFERENCES