

Optimal Control Problem and Power-Efficient Medical Image Processing Using Puma

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ABSTRACT: As a starting point of this paper we present a problem from mammographic image processing. We show how it can be formulated as an optimal control problem for PDEs and illustrate that it leads to penalty terms which are non-standard in the theory of optimal control of PDEs. To solve this control problem we use a generalization of the conditional gradient method which is especially suitable for non-convex problems. We apply this method to our control problem and illustrate that this method also covers the recently proposed method of surrogate functional from the theory of inverse problems. Graphics processing units (GPUs) are becoming an increasingly popular platform to run applications that require a high computation throughput. They are limited, however, by memory bandwidth and power and, as such, cannot always achieve their full potential. This paper presents the PUMA architecture - a domain-specific accelerator designed specifically for medical imaging applications, but with sufficient generality to make it programmable. The goal is to closely match the performance achieved by GPUs in this domain but at a fraction of the power consumption. The results are quite promising - PUMA achieves up to 2X the performance of a modern GPU architecture and has up to a 54X improved efficiency on a floating-point and memory-intensive MRI reconstruction algorithm.

KEYWORDS: generalized conditional gradient method, surrogate functional, image processing, optimal control of PDEs

I. INTRODUCTION

For many years medical imaging has aimed at developing fully automatic, software based diagnostic systems. However, the success of those automatic systems is rather limited and the human expert is as much responsible for the final diagnosis as in previous years. Hence, growing effort has been devoted to enhancing the techniques for presenting the medical images as well as additional information. In Germany a particular effort is made in mammography, i. e. X-ray scans of the female breast for early detection of breast cancer. The process of examination by the medical experts is divided into a very short recognition phase (< 1 sec.) and a second verification phase (≈ 1 min.). During the recognition phase, the expert first recognizes the coarse features, then more and more fine features. Tests have shown, that the experts usually form their decisions during this very short recognition phase. Nevertheless, the verification phase is the more critical one. The critical and difficult cases, where the recognition phase does not end with a preliminary diagnosis, most often applies to women in the early stages of cancer. During the verification phase the expert shifts forwards and backwards, thereby alternating in examining small details and in catching an overall impression of the location of critical patterns within the organ. The advent of programmable graphics processing units, or GPUs, for general-purpose computing is one of the major steps taken in computing over the last few years. These GPGPUs which, in the past, have been predominantly used for gaming and advanced image and video editing are now being used by many developers to accelerate inherently parallel programs in several other domains. Indeed, considerable amounts time and engineering effort are often spent in order to modify programs so that they may run effectively on GPUs. Several different application domains observe remarkable speedups when using GPUs, including the following [18]:

- 4X to 100X speedup on medical applications, such as biomedical image analysis, 3D reconstruction of tissue structures for large sets of microscopic images and accelerating MRI reconstructions.
- 8X to 260X speedup on electronic design automation, such as power grid analysis and statistical static timing analysis.
- 4X to 327X speedup on physics applications, such as weather prediction and astrophysics.
- 11X to 100X speed up financial applications such as instrument pricing using Monte-Carlo methods.

II. MOTIVATION FROM MEDICAL IMAGING

This process can be supported by presenting the expert different versions of the original image during close up and normal sub phases. More precisely, the expert sees a version with contrast enhanced small details in a close up phase ('fine scale'), while he sees an image which preserves all major edges but smoothes within regions ('coarse scale') during the normal phase. For enhancing fine details in mammography images a variety of algorithm have been proposed. Many of them are based on the wavelet transform due to its property of dividing an image into different scale representations; see for example [7] and references therein. In this work we deal with the development of an optimized presentation for one cycle of the verification phase. To put the problem in mathematical terms, we start with a given image y_0 assumed to be a function

defined on $\Omega := [0, 1]^2$. The fine scale and the coarse scale image are denoted y_f and y_c respectively. Under the natural assumption of finite energy images we model them as functions in $L^2(\Omega)$. The goal is, to produce a movie (i. e. a time dependent function) $y: [0, 1] \rightarrow L^2(\Omega)$, from the given images y_0, y_f and y_c such that the movie starts in y_0 , i. e. $y(0) = y_0$,

the movie sweeps to the fine scale image and to the coarse scale image, e. g. $y(t) \approx y_f$ for $t \in [0, .2]$ and $y(t) \approx y_c$ for $t \in [.6, .8]$, the movie sweeps in a “natural” way. An example for a mammography image, a fine scale, and coarse scale image is shown in Fig 1. As a first guess one could try to make a linear interpolation between the fine scale and the coarse scale representation. This method has one serious drawback: It does not take the scale sweep into account, i. e. all fine details are just faded in rather than developing one after another.

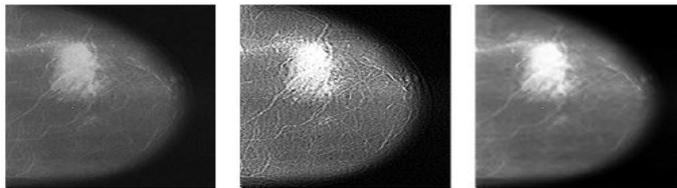


Fig 1: A mammography image. Left: original image y_0 , middle: fine scale image y_f , right: coarse scale image y_c
Modeling as an optimal control problem

III. PDES AND CONTROL PROBLEMS IN IMAGE PROCESSING

Parabolic partial differential equations are a widely used tool in image processing. Diffusion equations like the heat equation [14], the Perona-Malik equation [10] or anisotropic equations [13] are used for smoothing, denoising and edge enhancing. The smoothing of a given image $y_0 \in L^2(\Omega)$ with the heat equation is done by the solution of the equation

$$\begin{aligned} y_t - \Delta y &= 0 \text{ in } [0, 1] \times \Omega \\ y_\nu &= 0 \text{ on } [0, 1] \times \partial\Omega \\ y(0) &= y_0 \end{aligned} \tag{1}$$

Where y_ν stands for the normal derivative, i. e. we impose homogeneous Neumann boundary conditions.

The solution $y: [0, 1] \rightarrow L^2(\Omega)$ gives a movie which starts at the image y_0 and becomes smoother with time t . This evolution is also called scale space and is analyzed by the image processing community in detail since the 1980s. Especially the heat equation does not create new features with increasing time, see e. g. [5] and the references therein. Thus, the heat equation is well suited to model a sweep from a fine scale image y_f to a coarse scale image y_c . Hence, we take the image y_f as initial value. To make the movie y end at a certain coarse scale image y_c instead of its given endpoint $y(1)$ we propose the following optimal control problem:

$$\begin{aligned} \text{Minimize} \quad & J(y, u) = \frac{1}{2} \int_{\Omega} |y(1) - y_c|^2 dx + \frac{\alpha}{2} \int_0^1 \int_{\Omega} |u|^2 dx dt \\ \text{subject to} \quad & y_t - \Delta y = u \text{ in } [0, 1] \times \Omega \\ & y_\nu = 0 \text{ on } [0, 1] \times \partial\Omega \\ & y(0) = y_f. \end{aligned} \tag{2}$$

In other words, the diffusion process is forced to end in y_c with the help of a heat source u .

III.1: Adaption to image processing: The above described problem is classical in the theory of optimal control of PDEs, though not well adapted to image processing. The solution of this problem may have several drawbacks: The control u will be smooth due to the regularization and have a large support. This will result in very smooth changes in the image sequence y and, more worse, in global changes in the whole image. To overcome these difficulties, different norms can be used for regularization. A widely used choice in image processing is to use Besov norms because they are appropriate to model images. Besov norms can be defined in different ways, e. g. in terms of moduli of smoothness [12] or in terms of Littlewood-Paley decompositions [6]. Here we take another viewpoint and define the Besov spaces via norms of wavelet expansions [2,

9]. For a sufficient smooth wavelet ψ the Besov semi norm of a function f on a set $M \subset \mathbf{R}^d$ is defined as

$$|f|_{B_{p,q}^s(M)}^q = \sum_j \left(2^{sjp} 2^{j(p-2)d/2} \sum_{i,k} |\langle f, \psi_{i,j,k} \rangle|^p \right)^{q/p} \tag{3}$$

Where j is the scale index, k indicates translation and i stand for the directions. The Besov space $B_{p,q}^s(M)$ is defined as the functions $f \in L^p(M)$ that has a finite Besov semi norm. See [6, 9] for a more detailed introduction to wavelets and Besov spaces.

III.2: The solution of the PDE and the control-to-state mapping: The solution of the heat equation is a classical task. If we assume that the initial value y_f is in $L^2(\Omega)$ and the control u is in $L^2([0, 1] \times \Omega)$ the solution y is in $L^2(0, 1; H^1(\Omega)) \cap C([0, 1], L^2(\Omega))$. Especially y is continuous with respect to time and the point evaluation $y(1)$ makes sense, see e. g. [8]. In our

case the solution operator $u \mapsto y$ is affine linear, due to the nonzero initial value. We make the following modifications to come back to a linear problem: We split the solution into two parts. The non-controlled part y^n is the solution of

$$\begin{aligned} y_t^n - \Delta y^n &= 0 \\ y^n(0) &= y_f \end{aligned}$$

and the homogeneous part y^h is the solution of

$$\begin{aligned} y_t^h - \Delta y^h &= u \\ y^h(0) &= 0 \end{aligned} \tag{4}$$

(Both with homogeneous Neumann boundary conditions). Then the solution operator $G: u \mapsto y^h$ of equation (1) is linear and continuous from $L^2([0, 1], L^2(\Omega))$ to $L^2(0, 1, H^1(\Omega)) \cap C([0, 1], L^2(\Omega))$. With the help of the point evaluation operator we have the control-to-state mapping $K: u \mapsto y^h(1)$ linear and continuous from $L^2([0, 1], L^2(\Omega))$ to $L^2(\Omega)$. Then the solution is $y = y^n + y^h$ and we can focus on the control problem for y^h . Together with the thoughts of the previous subsection we end up with the following minimization problem:

Minimize

$$J(u) = \frac{1}{2} \|Ku - y_c + y^n(1)\|_{L^2(\Omega)}^2 + \alpha |u|_{B_{p,p}^s([0,1] \times \Omega)}^p \tag{5}$$

III.3. Solution of the optimal control problem: The minimization of the functional (2) is not straightforward. The non-quadratic constraint leads to a nonlinear normal equation which can not be solved explicitly. Here we use a generalization of the conditional gradient method for the minimization.

IV. THE GENERALIZED CONDITIONAL GRADIENT METHOD

The classical conditional gradient method deals with minimization problems of the form

$$\min_{u \in C} F(u), \tag{6}$$

here C is a bounded convex set and F is a possible non-linear function. One notices that this constrained problem can actually be written as an “unconstrained” one with the help of the indicator functional

$$I_C(u) = \begin{cases} 0 & u \in C \\ \infty & u \notin C \end{cases} .$$

With $\Phi = I_C$, problem (3) thus can be reformulated as

$$\min_{u \in H} F(u) + \Phi(u). \tag{7}$$

To illustrate the proposed generalization, we summarize the key properties of F and Φ : F is smooth while Φ may contain

non-differentiable parts. The minimization problem with Φ alone is considered to be solved easily while the minimization of

F is comparatively hard. The influence of Φ is rather small in comparison to F . With these assumptions in mind, the conditional gradient method can also be motivated as follows. Let $u \in H$ be given such that $\Phi(u) < \infty$. We like to find an

update direction by a linearized problem. Since Φ is not differentiable, we only linearize F :

$$\min_{v \in H} \langle F'(u) | v \rangle + \Phi(v). \tag{8}$$

The minimizer of this problem serves as an update direction. So this “generalized conditional gradient method” in the $(n+1)$ -

st step reads as follows: Let $u_n \in H$ be given such that $\Phi(u_n) < \infty$.

1. Determine the solution of (5) and denote it v_n .
2. Set s_n as a solution of

$$\min_{s \in [0,1]} F(u_n + s(v_n - u_n)) + \Phi(u_n + s(v_n - u_n)).$$

3. Let $u_{n+1} = u_n + s_n(v_n - u_n)$.

To ensure existence of a solution in Step 1 we state the following condition: Assumption 1 Let the functional $\Phi : H \rightarrow]-\infty, \infty]$ be proper, convex, lower semi-continuous and coercive with respect to the norm. Standard arguments from convex analysis yield the existence of a minimize in Step 1 of the algorithm [4]. So if F is G -ateaux-differentiable in H , the algorithm is well-defined. The convergence of the generalized conditional gradient method is analyzed in detail by the authors in [1]. The main result there is the following theorem.

Theorem 2 Let Φ satisfy Assumption 1 and let every set $E_t = \{u \in H \mid \Phi(u) \leq t\}$ be compact. Let F be continuously

Fréchet differentiable, let F_+ be coercive and u_0 be given such that $\Phi(u_0) < \infty$. Denote (u_n) thesequence generated by the generalized conditional gradient method. Then every convergent subsequence of (u_n) converges to a stationary point of $F +$

Φ . At least one such subsequence exists. Two remarks are in order: First, we notice that the theorem is also valid if the functional F is not convex. Second, the theorem only gives convergence to a stationary point which may seem unsatisfactory, specially if one wants to minimize non-convex functions. But this does not have to be a drawback, as we will see in the next section:

1. Application: Here we show the application of the above described methodology. Since the effects can be seen more clearly in artificial images, we will not use original images. The artificial images we used are shown in Fig 2.



Fig 2: Images used for illustration. Left: fine scale image, right: coarse scale image.

For illustration we use the values $p = 1$ and $s = 3/2 + \epsilon > 3/2$ in the minimization problem (2), since this is close to the BV-norm and we have $B_{1,1}^{3/2+\epsilon}([0, 1] \times \Omega) \subset L^2([0, 1] \times \Omega)$ compactly.

The results are presented in Figure 3. The figure shows a comparison of the linear interpolation, the pure result of the application of the heat equation and the result of the optimal control problem. One sees that the linear interpolation is only fading out the details. In the uncontrolled result (middle column) the details are vanishing one after another but the process does not end in the desired endpoint. The result of the optimal control problem (right column) exhibits both a nice vanishing of the details and end in the given endpoint.

V. THE ADVANTAGES OF GPUS

GPUs have many appealing hardware features. Firstly, they lend themselves very well to both thread-level and data-level parallelism. Thread-level parallelism (TLP) is exploited by having a large number of independent processing elements (PEs) on the GPU, each with its own set of functional units (FUs) and local storage. Individual threads can quite cleanly be assigned, either statically by the programmer or dynamically by the hardware, to each of these PEs and inter-thread communication is made possible by some form of interconnect fabric or through local storage such as caches. Programs with a large amount of data-level parallelism (DLP) can make use of vector-SIMD units in these PEs which allow a single instruction to perform an operation on several data at the same time. DLP can also be extracted in programs with compute-intensive loops that have little or no interiteration dependencies by executing operations from different iterations within a single SIMD instruction. Secondly, GDDR RAM and its increasingly fast successor's have allowed for GPUs to have access to an immense amount of memory bandwidth. The AMD Radeon HD 4870 - the first GPU to support GDDR5 memory - has a memory bandwidth of up to 115 GB/s. Above all, GPUs are commodity hardware products commonly available as a part of many desktop and laptop computers. The tools to program them are also easily available; NVIDIA's Compute Unified Device Architecture (CUDA) package, for example, is free to download from their website [15]. CUDA is a general purpose parallel computing architecture which consists of the CUDA instruction set and the compute engine in the GPU. It provides a small set of extensions to the C programming language, which enables straightforward implementation of parallel algorithms on the GPU. CUDA also supports scheduling the computation between CPU and GPU, such that serial portions of applications run on the CPU and parallel portions are mapped to the GPU. Individual cores in Intel's up-and-coming Larrabee processor

implement the ubiquitous x86 ISA [23], allowing users to use a host of already-existing development tools to port their applications to it. Server products like Tesla [17] with even more compute power are also available.

VI. THE QUEST FOR PROGRAMMABLE AND SPECIALIZED HARDWARE

A wide range of architectures, in addition to GPUs, have been designed before to address the problem of providing high performance computation efficiently. These solutions maintain or sacrifice programmability to various degrees depending on the domain they target. Fig 3 shows the performance (on the y-axis) and programmability (on the x-axis) expectations from various architecture styles. The numbers next to each of the ovals shows the approximate performance-power ratio offered by each of these solutions. General purpose processors (GPPs) which fall on the lower right corner of the figure, are highly programmable solutions but are limited in terms of the peak performance they can achieve. Further, structures like instruction decoders and caches that are needed to support programmability consume energy. This results in a very low computational efficiency of about 1 MIPS-per-mW, for example, for the Intel Pentium-M processor. On the other end of the spectrum are Application-specific Integrated Circuits (ASICs). ASICs are custom-designed specifically for a particular problem, without extraneous hardware structures. Thus, ASICs have a high computational density with hardwired control, resulting in high computation efficiency up to 1,000 to 10,000 times more than that of GPPs. The space between these two extremes is populated by different solutions that have varying degrees of programmability. Application specific instruction-set processors (ASIPs) are processors with custom extensions for a particular application or application domain. They can be quite efficient when running the applications for which they are designed, and they are also capable of running any other application, though with reduced efficiency. Examples include processors from Tensilica and ARC, transport triggered architectures [3] and custom-fit processors [9]. Domain loop accelerators are designed to execute computation intensive loops present in media and signal processing domains. Their design is close to that of VLIW processors, but with a much higher number of function units, and consequently, a higher peak performance. Very long instruction words in a control memory direct all FUs every cycle. However, domain loop accelerators (LAs) have less flexibility than GPPs because only highly computationally-intensive loops map well to them. Some examples of architectures in this design space are RSVP [1] and CGRAs [14]. Coarse-grain adaptable architectures have coarser-grain building blocks compared to FPGAs, but, like FPGAs, still maintain bit-level reconfigurability. The coarser reconfiguration granularity improves the computation efficiency of these solutions. However, non-standard tools are needed to map computations onto them and their success has been limited to the multimedia domain. PipeRench [10], RaPiD [6] are some examples of coarse-grain adaptable architectures.

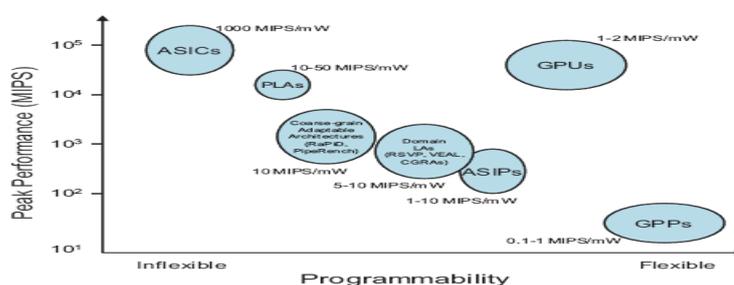


Fig. 3. Comparison of peak performance, power efficiency, and programmability of different architecture design styles.

1. **Programmable Loop Accelerators:** The programmable solutions shown in Figure 1 are all “universally” programmable, allowing any loop to be mapped on to them, although at varying degrees of efficiency. There is a wide gap between the efficiency that can be achieved by ASICs and the efficiency that can be achieved by these programmable solutions. There are, for example, instances where there is a narrow requirement of flexibility. Using any of these above solutions is overkill for these instances as these solutions sacrifice too much efficiency for the needed flexibility. Further, most of the middle ground solutions listed above do not offer any support for fast floating point computation, which limits the number of applications that they can be used for. This work advocates an open area in the design space where a non-trivial amount of programmability is provided in terms of intraprocessor communication, functionality and storage, but the application and domain-specific design, as a whole, resembles an ASIC more than a processor. The design point is labeled Programmable Loop Accelerator, or PLA (not to be confused with programmable logic array).

TABLE 1. Medical application characteristics

Benchmark	#instrs	%FP	Data Req'd <i>bytes</i> <i>instr</i>
MRI.FH	38	42	0.95
MRI.Q	34	35	1.06
CT.segment	26	42	1.38
CT.laplace	20	30	1.20
CT.gauss	22	32	1.09

VII. TARGETING MEDICAL APPLICATIONS

Medical imaging devices are generally large, bulky and expensive machines that have very limited portability and consume large amounts of power. There is an increasing focus on reducing the power of these medical imaging devices [20]. In order to address this issue, this work focuses on principle components of Computed Tomography

(CT) and Magnetic Resonance Imaging (MRI) image processing and reconstruction. A CT scan involves capturing a composite image from a series of X-Ray images taken from various angles around a subject. It produces a very large amount of data that can be manipulated using a variety of techniques to best arrive at a diagnosis. Oftentimes, this involves separating different layers of the captured image based on their radio densities. A common way of accomplishing this is by using a well known image-processing algorithm known as "image segmentation". In essence, image segmentation allows one to partition a given image into multiple regions based on any of a number of different criteria such as edges, colors, textures, etc. Being able to partition images in this manner allows for studying of isolated sections of the image rather than of all the information that was captured. The segmentation algorithm used in this work has three main floating-point-intensive components: Graph segmenting (CT.segment), Laplacian filtering (CT. Laplace) and Gaussian convolution (CT. gauss).

Laplacian filtering highlights portions of the image that exhibit a rapid change of intensity and is used in the segmentation algorithm for edge detection. Gaussian convolution is used to smooth textures in an image to allow for better partitioning of the image into different regions. An MRI scan, instead of using X-Rays, uses a strong magnetic and radio frequency fields to align, and alter the alignment of, hydrogen atoms in the body. The hydrogen atoms then produce a rotating magnetic field that can be detected by the MRI scanner and converted to an image. The main computational component of reconstructing an MRI image is calculating the value of two different vectors, known here as MRI.FH and MRI.Q, respectively (explained in more detail in [13], [24]). Table I shows some characteristics of the benchmarks in consideration. All of these benchmarks are floating-point-intensive and require large amounts of data for the computation they perform, especially when compared to the 0.15 bytes/instruction supported by the GTX 280 GPU mentioned earlier.

The main loops in these benchmarks are "do-all" loops - there are no inter-iteration dependences. Prior work in this field has predominantly focused on using commercial products to accelerate medical imaging. For instance, in [11], the authors port "large-scale, biomedical image analysis" applications to multi-core CPUs and GPUs, and compare different implementation strategies with each other. In [21], the authors study image registration and segmentation and accelerate those applications by using CUDA on a GPU. In [24], the authors use both the hardware parallelism and the special function units available on an NVIDIA GPU to dramatically improve the performance of an advanced MRI reconstruction algorithm.

There are several other such examples of novel work in this field. In contrast with such research, this work focuses on designing a new, highly efficient, microarchitecture and architecture with the specific hardware requirements of medical imaging in consideration.

VIII. PUMA

PUMA, *Parallel micro-architecture for Medical Applications*, is a tiled architecture as shown in Fig 4. It is specifically designed to maximize power-efficiency when executing medical imaging applications while still retaining full programmability. Each tile in PUMA is an instance of a specialized PLA - a generalized loop accelerator. The PLA tiles are connected to their neighboring tiles and to the external interface through a high-bandwidth mesh of on-chip routers.

1. **Background:** Fig. 5 shows the hardware schema for the single-function loop accelerator [7], [5]. The LA is designed to efficiently execute a modulo scheduled loop [19] in hardware. The lengths of the schedule, and the corresponding run-time of the loop, are determined by the *initiation interval* (Π) - the number of cycles between the beginnings of successive iterations of the loop. Thus, a lower Π corresponds to a shorter schedule and increased performance. The modulo schedule contains a *kernel* that repeats every Π cycles and may include operations from multiple loop iterations. The LA is designed to exploit the high degree of parallelism available in modulo scheduled loops with a large number of function units (FUs). Each FU performs a specific set of functions that is tailored for the particular loop. Each FU writes to a dedicated shift register file (SRF); in each cycle, the contents of the registers shift downwards to the next register. Point-to-point wires from the registers to FU inputs allow data transfer from producers directly to consumers. Multiple registers may be connected to each FU input; a multiplexer (MUX) is used to select the appropriate one. Since the operations executing in a modulo scheduled loop are periodic, the selector for this MUX is essentially a modulo counter. In addition, a central register file (CRF) holds static live-in register values that cannot be stored in the SRFs. The schema described is a template that is customized for the particular loop being accelerated. The number, types, and widths of the FUs, the widths and depths of the SRFs, and the connections from the SRFs to the FUs are all determined from the loop. During synthesis, the loop is first modulo scheduled to meet a given performance requirement, and then the details of the LA datapath are determined from the communication patterns in the scheduled loop. The control path for the single-function LA consists of a finite state machine with Π states corresponding to each of time slots in the kernel of the modulo schedule. In each state, control signals direct the execution of FUs (for FUs capable of multiple operations) and control the MUXes at the FU inputs. Finally, a Verilog HDL realization of the accelerator is generated by emitting modules with pre-defined behavioral Verilog descriptions that correspond to the datapath elements. A simulation environment is used to verify that the Verilog properly implements the loop. Finally, gatelevel synthesis, placement, routing, power analysis and post-synthesis verification are performed on the design.

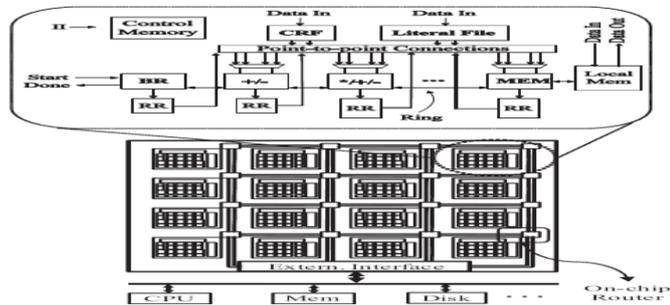


Fig. 4. PUMA. Each tile comprises of a programmable loop accelerator (template pictured) and the control and data memories required for its operation. On-chip routers transfer data between each tile and the external interface.

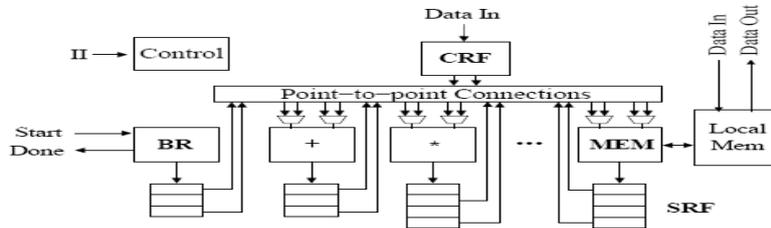


Fig. 5. Template for single-function loop accelerator.

2. PUMA Architecture

2.1. **Base line PLA Design:** A PLA is generalized loop accelerator, created by modifying the template datapath shown in Figure 5. A generic datapath template for the PLA is illustrated on the right side of Figure 4. The accelerator is designed for a specific loop at a specific throughput, but contains a more general datapath than the single-function LA to allow for different loops to be mapped onto the hardware [8]. These generalizations provide the LA with flexibility in functionality, storage, control and communication. To provide functionality, simple modifications were made to FUs in order for them to support more operations; adders (both integer and floating-point) are generalized to adder/subtractor units, left-shift units are generalized to left/right rotators, every FU can execute an identity operation to act like a move instruction, etc. Even load-store units can be generalized to integer adder/subtractor units if they already had the functionality to compute indirect addresses. Further, the input-muxes to FUs are redesigned to allow for operand-swapping as well. The SRFs used in the LA have limited addressability and fixed lifetimes for variables. To overcome these constraints and provide more generality, these SRFs are replaced with rotating-register files (RRs). To improve controllability, the LA's finite state machine is replaced with a control memory, the contents of which can be changed based on the loop that needs to be executed. Further, numerical constants which were hard-coded in the LA are instead stored in a literal register file. To generalize communication, the PLA has a bus (in addition to the point-to-point connections) that connects all the RRs to all the FUs. To reduce the hardware cost of having a potentially long bus, its width is limited to one operand, but has a predictable latency of one cycle.

$$\begin{aligned}
 &\text{Maximize:} \\
 &\sum_{i \in T_\alpha} \sum_{j \in T_\beta} C_{ij} \quad \forall \alpha \neq \beta \\
 &\text{Subject to:} \\
 &(1) \quad \sum_{j=0}^{\#FUs} X_{ij} = 5 \quad i \in [0, \#FUs) \\
 &(2) \quad X_{ii} = 1 \quad i \in [0, \#FUs) \\
 &(3) \quad C_{ii} = 1 \quad i \in [0, \#FUs) \\
 &(4) \quad X_{ij} = X_{ji} \quad i, j \in [0, \#FUs) \\
 &(5) \quad C_{ij} = C_{ji} \quad i, j \in [0, \#FUs) \\
 &(6) \quad C_{ij} \leq X_{ij} + I_{ij} \quad i, j \in [0, \#FUs)
 \end{aligned}$$

Fig. 6. ILP formulation for FU arrangement on the PUMA ring

2.2. **PUMA PLA:** The PLA bus is not always a viable solution. One main disadvantage with the bus is that it is not a scalable solution for larger PLAs with many FUs. Further, the bus only carries a single operand per cycle, limiting the amount of programmability available in the PLA and the sequences of opcodes that can be executed in parallel. To overcome these limitations, the intra-PLA communication fabric in PUMA is changed to a ring. A ring allows for as many operands to be transferred as there are connections to FUs. It does have its limitations, however. The assumed single-cycle latency to transfer data between two arbitrary points in the PLA using the bus is no longer valid, as it takes one cycle to transfer an operand from one ring connection (or ring stop) to another. Also, the longer the distance an operand needs to travel on the ring, the more FUs that have to execute move instructions to propagate the operand along at each ring stop. These added instructions can potentially increase the loop's schedule length and reduce the accelerator's performance. In PUMA, the ring architecture actually consists of six rings (three sets of two rings going in opposite directions). The first set of rings has a

Bus/FU connector (or ring-stop) at every single FU. The second set of rings has a ring-stop at all the odd-numbered FUs, and the third set of rings has a ring-stop at all the even-numbered FUs. This effectively connects an FU/RF pair to its two neighbors and also to its neighbors' neighbors; i.e. every FU can communicate with itself or with other FUs one or two positions on either side of it on the ring. With this configuration, the number of cycles required to transmit data between any

two arbitrary FUs is no more than $\lceil \frac{\#FUs}{4} \rceil$, and regardless of the ordering of FUs on the ring, every possible producer-consumer pairing can be executed, provided sufficient time. In order to best maintain generality, we chose to arrange the FUs along the ring to allow maximum connectivity and to distribute the various types of FUs as evenly as possible. This was done by formulating the problem as an integer linear program (ILP) as shown in Fig 6. In the objective function, T_i and T_j are two different sets of FUs, each set having all and only the FUs of a particular type. The subscripts i and j are FU IDs and C_{ij} is a binary variable that is 1 if a connection exists between FU i and FU j . Essentially, this objective function aims to maximize the number of connections between different types of FUs, subject to the following constraints: In constraint set (1), X_{ij} is a binary variable that is 1 if FU i is "positioned" adjacent to FU j , implying that they are connected by the ring. Every FU, therefore, is "positioned" next to 5 other FUs: itself, its two neighbors and the two additional FUs neighboring its neighbors. Constraint sets (2) and (3) specify that every FU is "positioned" next to and connected to itself. Constraint sets (4) and (5) specify that all added connections are bidirectional. In constraint set (6), I_{ij} is a binary number that is 1 if a connection between FU i and FU j has already been inserted by the synthesis tool. This constraint enforces the rule that a connection between FU i and FU j can only exist if they are either "positioned" next to each other or are already connected. A 7th set of constraints was initially used which specified that there must always be a path between any two FUs with exactly $\lceil \frac{\#FUs}{4} \rceil$ connections between them. This constraint was used to prevent insular sets of 5 FUs or sets of 5 FUs connected linearly rather than in a ring (i.e. without a direct connection between the two ends). While this problem might occur in theory, the preexisting connections put in place by the synthesis system prevent it from happening in practice and these constraints were removed to reduce the size of the ILP. Once the optimal solution is obtained, the values of the X_{ij} variables provide a unique ring arrangement.

IX. EXPERIMENTS AND RESULTS

1. **Setup:** All the PLAs in this work were synthesized for (and run at) a frequency of 450 MHz. The logic synthesis was done using Synopsys Design Compiler 2006-06 and Synopsys Physical Compiler 2006-06, targeting a 65nm process technology with a nominal supply voltage of 0.9 Volts. Energy numbers were obtained using Synopsys PrimeTime-PX 2006-12. For the purposes of this study, we assume that a peak memory bandwidth of 142 GB/s is available to each PUMA system. This is the same amount of bandwidth afforded to the NVIDIA GTX280 processor.

2. **PLA Characteristics:** PUMA systems were built using PLAs for each of the five benchmarks in consideration (five systems, each composed entirely of multiple tiles of a single type of PLA). Table II shows various characteristics of these accelerators. The "Peak Perf." columns show

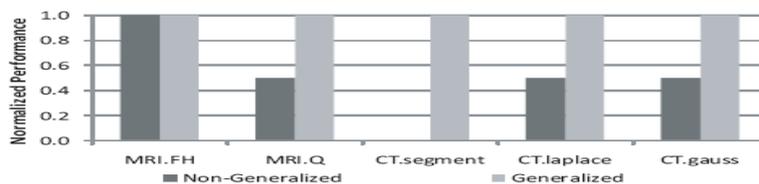


Fig. 7. Normalized performance of benchmarks on LA and PUMA PLA designed for MRI.FH

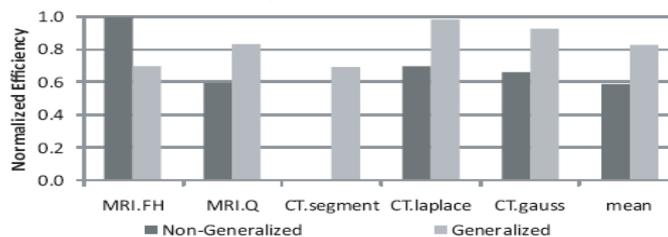


Fig. 8. Normalized efficiency of benchmarks relative to MRI.FH

The throughput when executing floating-point operations and integer operations, respectively, in billions of operations per second. The next column shows the minimum bandwidth required by each application to prevent starvation. Finally, the last column shows the total number of tiles of each PLA that would be present in a PUMA system. The number of tiles was chosen to prevent data starvation, to make the most efficient use of the resources available. For example, the number

of tiles in a system with MRI.FH tiles is $\lceil \frac{142}{16.2} \rceil$ or 9. Fig 8 shows the normalized performance difference between the non-generalized and generalized loop accelerators across various

Benchmarks, to illustrate the effects of the modifications made to the baseline accelerator to increase programmability. Each of the different benchmarks were compiled for the MRI.FH accelerator. The left column for each benchmark shows its normalized performance. The benchmarks MRI.Q, CT. Laplace and CT. gauss suffered a 50% reduction in performance; i.e. their II values had to be doubled, from 1 to 2, in order for them to execute on the baseline loop accelerator. The benchmark CT.segment could not be compiled for the MRI.FH accelerator at all. For each benchmark, the column on the right shows the achieved performance on the generalized accelerator, with the hardware modifications specified in section III-B1. As shown, these modifications allowed all the benchmarks to run at full performance, at minimum II. Fig 8 shows a graph similar to that in Fig 7, but shows the normalized efficiency in terms of the accelerator's performance to-power ratio. Due to the increase in overall performance provided by the generalizations, the benchmarks MRI.Q, CT.laplace and CT.gauss had a significant increase in efficiency despite the power overhead of the additions. The MRI.FH benchmark, however, which would not experience any improved performance from the generalizations loses efficiency due to the increase in the accelerator's power consumption. On average, the generalizations increased the accelerator's efficiency increased by approximately 40%.

3. Commodity GPGPU Comparison: While other architectures may certainly be used for this domain, GPGPUs are the solutions that are currently in use in many medical imaging applications and, therefore, the most suitable comparison point. For this reason, we assessed the performance and efficiency of five NVIDIA GPUs.

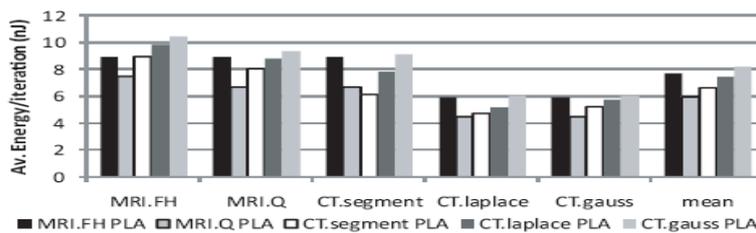


Fig. 9. Average energy consumed (per iteration) by each benchmark while running on PUMA systems designed around different PLAs

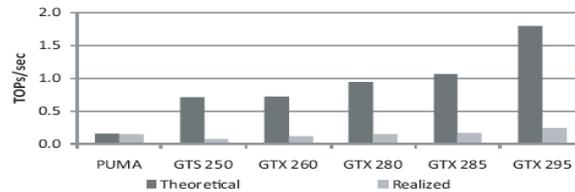


Fig. 10. Achieved performance of the MRI.FH benchmark (in trillions of operations) on the MRI.FH PUMA system and on various NVIDIA GPUs based on the GT200 architecture

Fig9 shows the result of direct performance comparisons between an MRI.FH PUMA system and the GPUs in consideration. The column on the left shows the total compute capability of each of the processors. The column on the right shows the realized performance while executing the MRI.FH benchmark, accounting for bandwidth restrictions. PUMA achieves a very small fraction of the peak performance offered by the GPUs, between 8.6% of the dual-GPU GTX 295 and 21.8% of the GTS 250. This gap changes dramatically, however, when accounting for the bandwidth-intensive nature of the application in question. PUMA delivers between 63% (on the dual-GPU GTX 295) and 2X the performance (on the GTS 250) of the GPUs. The case for PUMA is further underscored by examining the GPUs' power efficiency, as shown in Fig 10.

This graph shows how many times more efficient, in terms of number of operations per Watt, PUMA systems are relative to the GPUs in consideration. These values range from 20X, for the most complex benchmark running on the most efficient GPU, to 54X, for the least complex benchmark running on the least efficient GPU.

X. CONCLUSION

We have seen that the application of the theory of optimal control of PDEs to image processing problems is a fruitful field of research. Besides promising result, even for easy models like the linear heat equation, new interesting mathematical problems arise, like the treatment of non-quadratic penalty terms. For future research, better adapted PDEs (like the anisotropic diffusion equations) could be investigated. The PUMA architecture is a power-efficient accelerator system designed specifically for efficient medical image reconstruction. It consists of tiles of programmable loop accelerators - ASICs with added hardware to support general-purpose computing - designed around the computation requirements of the image reconstruction domain. As applications in this domain are normally executed on very high-performance GPGPUs, the latest NVIDIA GPU architecture was used to gauge the performance and efficiency of PUMA. The results are very encouraging - PUMA achieves up to 2 times the performance of a modern GPU architecture and has up to 54 times the power efficiency.

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