Dynamic Reuse of Memory in 2D Convolution Applied to Image Processing

Martin Casabella, Sergio Sulca, Ivan Vignolles, Ariel L. Pola

Escuela de Ingeniería en Computación
Facultad de Ciencias Exactas Físicas y Naturales
Universidad Nacional de Córdoba
Email: martin.casabella@gmail.com, ser.0090@gmail.com, ivignolles@alumnos.unc.edu.ar, arielpola@gmail.com

Abstract—**En este artículo se presenta una arquitectura de hardware para realizar una convolución 2D en una FPGA cuando no se puede instanciar suficiente memoria RAM para poder alojar la imagen completa. Se priorizó la velocidad de procesamiento, el uso eficiente de los recursos y un diseño escalable donde se pudieran agregar tantas operaciones de convolución en paralelo como se desee sin deber hacer grandes modificaciones en el diseño.**

1. Introduction

The origin of digital image processing, due to the high level of processing they require, is directly related to the development and evolution of computers. Most filters for images that focus, blur, enhance edges and detect edges, among others, use convolution as a mathematical operation. Image processing is a very extensive research area with a large number of applications in multiple fields such as medicine, engineering, navigation, aeronautics, among others. Applications recently, use convolutional neural networks (CNN) [1] where a fast and efficient convolution implementation is important.

The implementation of high speed image processing systems in FPGA has been a very active field. This is mainly due to the ability to take advantage of bit level parallelism, pixel level, neighborhood level and task level to accelerate the calculation. In addition, FPGAs are reconfigurable, allowing the flexibility that is often desired in neural networks. This coupling of processing, parallelism and flexibility at very high speed is what makes FPGAs a platform of choice for the development of these areas [2].

There are many examples in the literature of 2D convolution implementations, where most of the work focuses on high performance, resource reduction and the FPGA area. As shown in [3], BRAM memory blocks are one of the most used elements and consume more power. Therefore, the architectures seek to reduce the use of BRAMs by dividing them into two groups that consider the total or partial storage of the image. The first of them increases processing by applying parallelization as well as reusing resources to reduce complexity [4], [5]. On the other hand, architectures with partial storage partition the image into as many parts as convolders are used [6], [7]. The difficulty presented by these schemes is the non-modularization of

processing, making it difficult to increase the work rate by increasing parallelism.

In this document, we propose a high-performance, parallelizable hardware architecture and dynamic memory reuse with a linear increase in BRAM resources.

2. Image Processing Algorithms

Filtering is widely used in many applications, such as aeronautics, navigation, CNN, etc. They are applied as preprocessing to eliminate useless details and noise. One of the preferred devices for the development of these techniques is the FPGA, since it presents an excellent relationship between speed of development and flexibility in the implementation. It allows the parallelization of the process and the reconfiguration of the design in a very short time. We will focus on the efficient use of FPGA resources by applying the 2D convolution technique and reducing the dynamic range of the pixels in the image.

2.1. Convolution Operation

Bidimensional convolution is defined mathematically as

$$S(x,y) = \sum_{i=0}^{m-1} \sum_{j=0}^{n-1} K(i,j)I(x-i,y-j) \qquad (1)$$

where $I(x,y)$ is an image of size $(m \times n)$ pixels, $K(i,j)$ is a set of coefficients called Kernel of size $(h \times h)$ and $S(x,y)$ is convolution result of size $(m-2 \times n-2)$ pixels because we are considered valid convolution [8].

2.2. Change of Dinamic Range

We will focus on grayscale images because of their simplicity of implementation, but the design can be extrapolated to any other type of image. The pixels of the grayscale images $I(x,y)$ have a dynamic range between $[0, 255]$ and Kernel values $K(x,y)$ can be negative or positive. This work is part of a Deep Learning project where it is usual to operate with a dynamic range centered on zero. Therefore, we apply dynamic range expansion [9] and maximum norm [10] to rearrange $\mathcal{T}[I(x,y)] = I'(x,y)$ between $[0, 1]$ and

$\mathcal{T}[K(x,y)] = K'(x,y)$ between $[-1,1]$, respetively. Therefore, replacing in (1)

$$G'(x,y) = \sum_{i=0}^{m-1} \sum_{j=0}^{n-1} K'(i,j)I'(x-i,y-j). \qquad (2)$$

Finally, we apply the change of dynamic range on the output $\mathcal{T}[G'(x,y)] = G(x,y)$ again and obtain range between $[0,255]$. The complete process is detailed in Fig. X.

3. Architecture Design

3.1. Workflow

In this section, the design of the architecture is described in detail. The entire process is divided into three stages, pre-processing, processing and post-processing. The pre-processing consists of capturing the image and applying the detailed transformation in Section 2 using a Python script. In addition, the script divides the image into batch and is sent by the UART port to the FPGA. In the FPGA a microprocessor receives the information and communicates it to the convolution module using a 32 bits GPIO port. This port writes the BRAMs that store the data that will be filtered. Due to the limitations of the size of the FPGA, no other interface was used, such as Ethernet or DMA to read the image directly from a storage device. However, the proposed architecture is indifferent to these methods.

3.2. Convolution Module

A simplified architecture of the module is shown in Fig. 1, the **control unit** is the one in charge of handling the communication with the processor, in the **convolution unit** is where the sum of the products of the kernel's coefficients with the pixels happen, the **address generation unit (AGU)** manages the memory addresses, the **memory management unit (MMU)** manages how the values in memory must be read and written, and finally the **storage**, which is implemented with a set of columns of the FPGA's block RAM.

3.2.1. States. It is possible to see that during the work cycle, the convolution module goes through several states, as shown in Fig.2.

At the beginning the module must be in a **reset** state, waiting to be configured, to do so it must receive a reset signal.

State transition are controlled by coded instructions embedded in the 32 bits input frame.

The module will stay in the **reset** state until it receives the instruction to go to the **KernelLoad** state, in which state the kernel coefficients are loaded into the module. Then it goes to the **SizeLoad** state where the image's height is loaded, the need for this is explained in section 3.2.2. Once the module has gone through this setup states it will not come back to them until the entire image is processed. Now the processing loop start, first the module goes to the
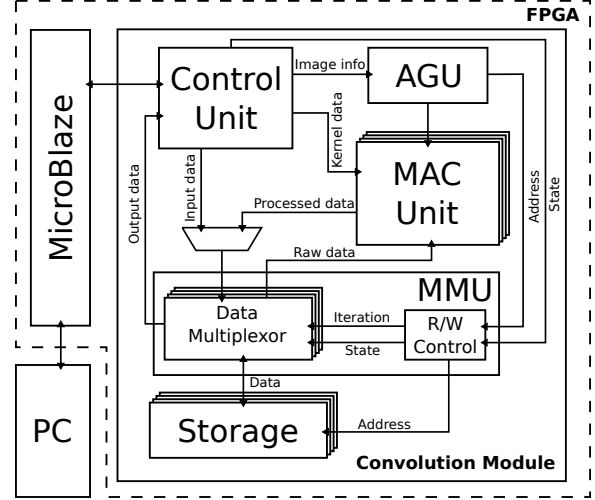

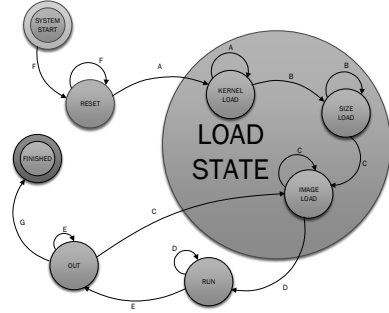Fig. 1. Simplified block diagram of the convolution module.

Fig. 2. Module's state diagram. A - Kernel load, B - Size load, C - Image load, D - Proccesing, E - Output, F - Reset

ImageLoad state and saves in the memory the batch, then is goes to the **Run** state where the processing occurs, while in processing the module can not be interrupted, i.e. it will ignore all instructions till the batch is processed, as soon as the processing is ended a notification will be emitted through Control Unit's *EoP* pin and the module will wait the instruction to got to the **Out** state where the already processed batch is send back to the microprocessor.

All this states logic is managed by the control unit, which depending on the state will communicate the corresponding control signal to the other components of the module.

3.2.2. Data storage. The kernel coefficients are stored in registers inside each convolution unit. In order to store a batch, the proposed approach is to arrange the block RAM memory in columns, where each of them will store a image's pixel column. Therefore a batch size is given by the height of the image and the number of memory columns, also the maximum height of the image will be restricted by the number of entries that each memory columns has.

During a batch processing, every pixel is read only once, so in order to do a more efficient use of the memory, the

pixels that were already used are overwritten with processed pixels. This approach reduces drastically the amount of memory needed because it reuses the same memory to store the input batch and the processed batch. In section 3.2.3 additional aspects of the design that reduces even more the memory use are explained.

3.2.3. Data processing. Given a $k \times k$ kernel, each convolution unit takes k adjacent memory columns as input. To produce the first processed pixel, a convolution unit first needs to load k pixels from each columns in such a way as to have $k \times k$ pixels loaded into it, then proceeds to multiply the pixels with the kernel's coefficients and sum everything. Finally, it truncates the result to reduce the bit length and the result is saved into the first position of the memory.

Once a pixel is processed, it is saved into the memory while the convolution unit shifts its image register, discarding the oldest k pixels and loading new k pixels, i.e. one from each input column, like a FIFO structure. This procedure is equivalent to shit the kernel vertically on the image. All this steps are synchronized in such a way that one processed pixel is generated in every clock cycle.

The AGU manages the memory's read and write address, it takes into account the latency between the clock cycle where the first pixel in the memory is read and the clock cycle where the first processed pixel is written in memory. This latency is equal to k plus some more clock cycles needed to latch values during the convolution, and the difference between the read address and the write address must be equal to it.

Until this point the analysis was over a single convolution unit, now the approach to work with multiple convolution units working in parallel is described.

As mentioned above, k columns are needed as input to produce one processed column in a convolution unit, that means that $2k$ columns are needed for two convolution units, however given the nature of the convolution operation, to produce a contiguous column it is necessary to shift the input columns by one. Hence, there is an overlap between the two inputs and this produces that even that k input columns are still needed per convolution unit, only $k+1$ different input columns are needed for all the units. For that reason, it was decided that multiple convolution units in parallel will produce contiguous processed columns, sharing the common input columns. With this approach, for N convolution units the number of memory columns needed gets reduced from $N \cdot k$ to $N+k-1$. That means that adding a new convolution unit only adds one new memory columns as Fig. 3 shows.

The same overlap explained above produces also that repeated data will be needed between one batch and the next one. In order to eliminate the need to send data that already is in the memory, some data is reused between iterations. Given N convolution units and a $k \times k$ kernel, a $N+k-1$ batch width is needed, but the processed batch has a width of N columns, i.e. one for each convolution unit, therefore the last $k-1$ memory columns are not overwritten and maintain the input data. This $k-1$ columns will be reused as the first columns of the next batch, thus the batch width
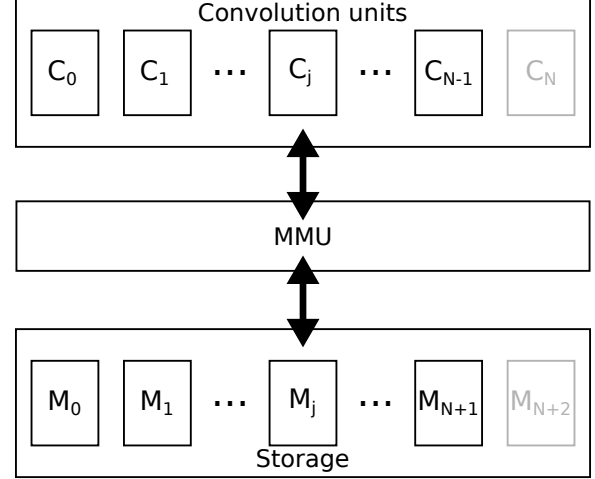

Fig. 3. Using shared inputs approach, only one new memory column is added for every new instantiated convolution unit.
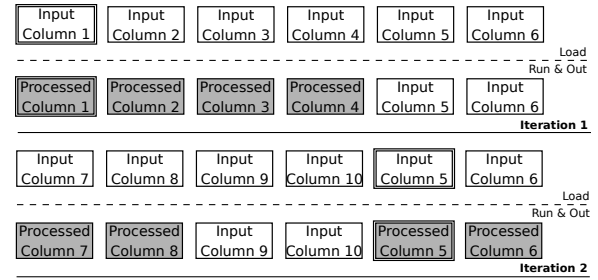

Fig. 4. Memory writing process using circular shift for $N = 4$ and $k = 3$.

is reduced to N, with the exception of the first batch that will still have a $N+k-1$ width.

A consequence of reusing the memory columns written by the previous batch is that in every iteration the position of the memory columns associated with each convolution unit describes a circular shift by N places. From the above it follows that a periodicity in the relation between the memory columns and the convolution units' inputs must exist, where the period It is the number of iterations necessary to get to the original memory columns - convolution units inputs relation, i.e. when $It(k-1)$ is a multiple of $N+k-1$. That is, there must be an integer m such that

$$\frac{It}{m} = \frac{N}{k-1} + 1 \qquad (3)$$

Figure 4 shows how the memories are written using this logic.

The logic is implemented in the MMU, which serves as an interface between the memories and the rest of the components, keeping them independent from the parallelism degree and the iteration number.

The MMU keeps track of the memory positions where it must save the input batch, the information which must feed to every convolution unit, the memory position where the processed data must be stored and the order in which the
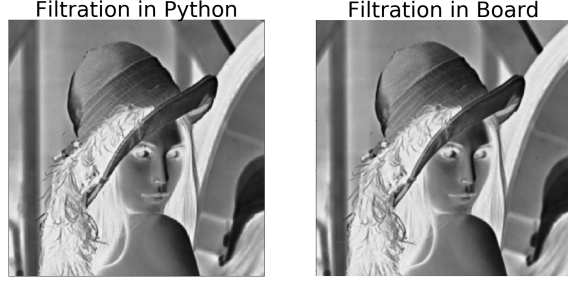
Fig. 6. On the left, an image processed in a PC using python. On the right the same image processed with the FPGA module.
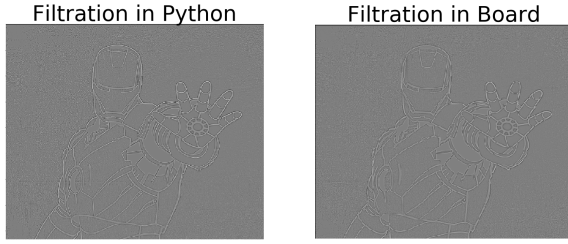

Fig. 7. On the left, an image processed in a PC using python. On the right the same image processed with the FPGA module.

Fig. 8. Normalized curve for BRAM resources utilization.

processed data must be returned, for each iteration. To do so, it has a finite state machine where each state corresponds to a set of positions named above. The number of states is given by eq. 3.

To route the incoming and outgoing data, the MMU has a set of multiplexers whose select lines are managed by the finite state machine (Data multiplexer and FSM in Fig. 1).

A comparison between the naive (which would be assign k independent memory columns to each convolution unit), the shared inputs and the shared inputs with circular shift approaches is shown in Fig. 5. Figure 5a shows the amount of memory required in function of the parallelism degree while Fig. 5b shows the amount of transferred bytes (considering that every byte transmitted is stored in only one memory register) for processing a 1600×1024 image and a 3×3 kernel. As shown in Fig. 5b, with the naive approach the transferred data is almost 3 times the image size (because a 3×3 kernel) and it does not get affected by the parallelism degree. The shared inputs approach tends to diminish the redundant data transferred as N increases, that is because the redundant data in a $N + k - 1$ batch is $k - 1$ and $\frac{k-1}{N+k-1}$ tends to zero as N increases. The shared inputs with circular shift approach is the best case scenario where no redundant data is transferred at all.

4. Implementation and Results

To ensure the proper functioning of the architecture a negative filter was coded in python and applied to an image, then the same filter was applied using the FPGA module. The results are showed in Fig. 6.
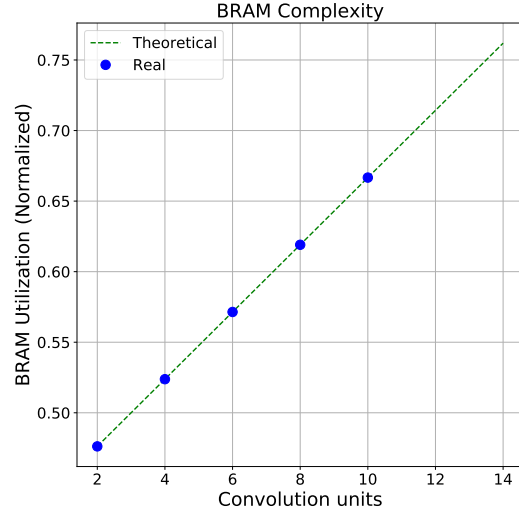
The design was implemented in an Arty 7 FPGA from Xilinx. The architecture was synthesized for different number of convolution units. The results obtained for BRAM utilization from the synthesis are compared with the estimated results in Fig. 8. As expected, the increase has a linear behaviour.

5. Conclusion

Acknowledgments

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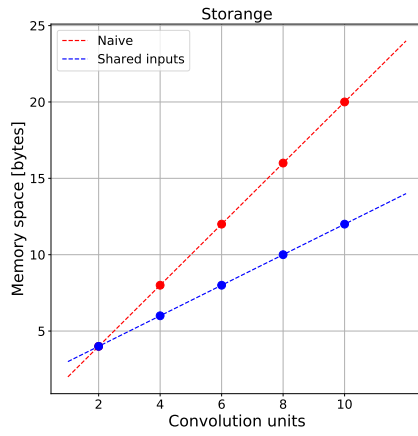
(a) Amount of memory required.
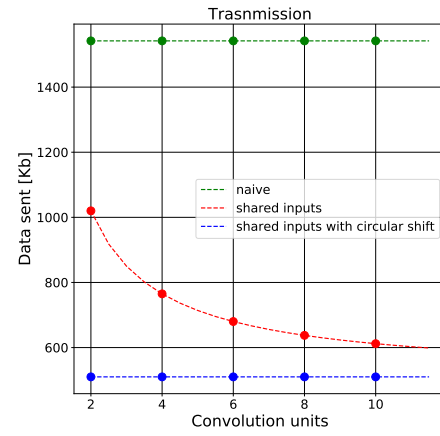
(b) Amount of data transmitted for an image of 1600×1024px and a 3×3 kernel.

Fig. 5. Comparison between different approaches.

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