

# Spiking Neural P system without delay simulator implementation using GPGPUs

Francis Cabarle

Algorithms and Complexity Laboratory  
Department of Computer Science  
University of the Philippines Diliman  
fccabarle@up.edu.ph

Henry Adorna

Algorithms and Complexity Laboratory  
Department of Computer Science  
University of the Philippines Diliman  
hnadorna@dcs.upd.edu.ph

## ABSTRACT

This paper presents a parallel simulator for a type of P system known as spiking neural P system (SNP system) using general purpose graphics processing units (GPGPUs). GPGPUs, unlike the more conventional and general purpose, multi-core CPUs, are used for parallelizable problems due to their architectural optimization for parallel computations.

Membrane computing or P systems on the other hand, are cell-inspired computational models which compute in a maximally parallel and non-deterministic manner. SNP systems, w/c compute via time separated spikes and whose inspiration was taken from the way neurons operate in living organisms, have been represented as matrices.

The matrix representation of SNP systems provides a crucial step into their simulation on parallel devices such as GPGPUs. Simulating the highly parallel nature of SNP systems necessitates the use of hardware intended for parallel computations. The simulator algorithms, design considerations, and implementation are presented. Finally, simulation results, observations, and analyses using an SNP system that generates all numbers in  $\{N - 1\}$  are discussed.

## Keywords

Membrane computing, Parallel computing, GPU computing

## 1. INTRODUCTION

### 1.1 Parallel computing: Via graphics processing units (GPUs)

The trend for massively parallel computation is moving from the more common multi-core CPUs towards GPGPUs for several significant reasons [7][8]. One important reason for such a trend in recent years include the low consumption in terms of power of GPGPUs compared to setting up machines and infrastructure which will utilize multiple CPUs in

order to obtain the same level of parallelization and performance[9]. Another more important reason is that GPGPUs are architected for massively parallel computations since unlike the architectures of most general purpose CPUs, a large part of GPGPUs are devoted for arithmetic operations and not on control and caching [7][8]. Arithmetic operations are at the heart of many basic operations as well as scientific computations, and these are performed with larger speedups when done in parallel, by GPGPUs over CPUs.

### 1.2 Parallel computing: Via Membranes

Membrane computing or its more specific counterpart, a P system, are Turing complete computing models that perform computations nondeterministically, exhausting all possible computations at any given time. This type of unconventional model of computation was introduced by Gheorghe Păun in 1998 and takes inspiration, similar to other members of natural computing, from nature[4][5]. Specifically, P systems try to mimic the constitution and dynamics of the living cell: the multitude of elements inside it, and their interactions within themselves and their environment, or outside the cell's skin membrane.

SN P systems differ from other types of P systems precisely because they are mono-membranar and only use one type of object in its computation. These characteristics, among others, are meant to capture the workings of a special type of cell known as the neuron. Neurons, such as those in the human brain, communicate or 'compute' by sending indistinct electro-chemical signals more commonly known as spikes. Information is then communicated and encoded not by the spikes themselves, since the spikes are unrecognizable from one another, by means of time duration as well as the number of spikes sent/received from one neuron to another, oftentimes under a certain time interval[1]. The time duration between two spikes, or several successive spikes, transmit information from one cell to another.

It has been shown that SN P systems, given their nature, are representable by matrices[2][3]. This representation allows design and implementation of an SN P system simulator using parallel computing machines such as GPGPUs.

### 1.3 Simulating SNP systems in GPGPUs

Matrix operations and their algorithms have been studied and efficiently implemented in GPGPUs [10][11]. Thus the matrix representation of SNP systems bridges the gap between the highly theoretical yet still computationally pow-

erful SNP systems and the applicative and more tangible GPGPUs, via an SNP system simulator. The design of the simulator, including the algorithms devised, architectural considerations, are then implemented using a particular type of GPGPU, namely NVIDIA CUDA (compute unified device architecture). NVIDIA CUDA extends the widely known ANSI C programming language and makes parallel computations, via GPGPUs manufactured by NVIDIA.

This paper starts out by introducing and defining the type of SNP system that will be simulated. Afterwards the NVIDIA CUDA model and architecture are discussed, baring the scalability and parallelization CUDA offers. Next, the design of the simulator, constraints and considerations, as well as the details of the algorithms used to realize the SNP system are discussed. The simulation results are presented next, as well as observations and analysis of these results. The paper ends by providing the conclusions and future work.

## 2. SPIKING NEURAL P SYSTEMS

### 2.1 Computing with SNP systems

The type of SNP systems focused on by this paper are those without delays i.e. those that spike or transmit signals the moment they are able to do so [2][3]. A variant, which allows for delays before a neuron produces a spike, are also available [1]. An SNP system without delay is of the form:

$$\Pi = (O, \sigma_1, \dots, \sigma_m, syn, in, out),$$

where:

1.  $O = \{a\}$  is the alphabet made up of only one object, the system spike  $a$ .
2.  $\sigma_1, \dots, \sigma_m$  are  $m$  number of neurons of the form

$$\sigma_i = (n_i, R_i), 1 \leq i \leq m,$$

where:

- a)  $n_i \geq 0$  gives the initial number of  $a$ s i.e. spikes contained in neuron  $\sigma_i$
- b)  $R_i$  is a finite set of rules of with two forms:
  - (b-1)  $E/a^c \rightarrow a$ , are known as *Spiking rules*, where  $E$  is a regular expression over  $a$ , and  $c \geq 1$ , such that  $c \geq 1$ .
  - (b-2)  $a^s \rightarrow \lambda$ , are known as *Forgetting rules*, for  $s \geq 1$ , such that for each rule  $E/a^c \rightarrow a$  of type (b-1) from  $R_i$ ,  $a^s \notin L(E)$ .
3.  $syn = \{(i, j) | 1 \leq i, j \leq m, i \neq j\}$  are the synapses i.e. connection between neurons.
4.  $in, out \in \{1, 2, \dots, m\}$  are the input and output neurons, respectively.

Furthermore, rules of type (b-1) are applied if  $\sigma_i$  contains  $k$  spikes,  $a^k \in L(E)$  and  $k \geq c$ . Using this type of rule uses up or consumes  $k$  spikes from the neuron, producing a spike to each of the neurons connected to it via a forward pointing arrow i.e. away from the neuron. In this manner, for rules of type (b-2) if  $\sigma_i$  contains  $s$  spikes, then  $s$  spikes are forgotten or removed once the rule is used. Rules of type (b-1) can be simplified with the notation

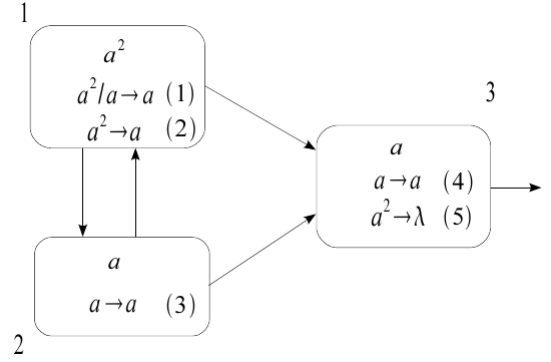


Figure 1: An SNP P system  $\Pi$ , generating all numbers in  $\{\mathbb{N} - 1\}$ , from [3].

$$(b-3) a^k \rightarrow a$$

where the regular expression  $E = a^k$ , again consuming  $k$  spikes and producing a spike.

The non-determinism of SNP systems comes with the fact that more than one rule of the several types are applicable at a given time, given enough spikes. The rule to be used is chosen non-deterministically in the neuron. However, only one rule can be applied or used at a given time [1][2][3]. The neurons in an SNP system operate in parallel and in unison, under a global clock [1]. For Figure 1 no input neuron is present, but neuron 3 is the output neuron, hence the arrow pointing towards the environment, outside the SNP system. The SNP system in Figure 1 is a 3 neuron system whose neurons are labeled (neuron/ $\sigma_1$  to 3) and whose rules have a total system ordering from (1) to (5)

### 2.2 Matrix representation of SNP systems

A matrix representation of an SNP system makes use of the following vectors and matrix definitions [2][3]. It is important to note that, just as in Figure 1, a total ordering of rules is in order.

*Configuration vector*  $C_k$  is the vector containing all spikes in every neuron on the  $k$ th computation step/time, where  $C_0$  is the initial vector containing all spikes in the system at the beginning of the computation. For  $\Pi$  (in Figure 1) the initial configuration vector is  $C_0 = \langle 2, 1, 1 \rangle$ .

*Spiking vector* which shows, at a given configuration  $C_k$ , if a rule is applicable (has value 1) or not (has value 0 instead). For  $\Pi$  we have the spiking vector  $S_k = \langle 1, 0, 1, 1, 0 \rangle$  given  $C_0$ . Note that a 2nd spiking vector,  $S_k = \langle 1, 0, 1, 1, 0 \rangle$ , is possible if we use rule (2) over rule (1) instead (but not both at the same time, hence we cannot have a vector equal to  $\langle 1, 1, 1, 1, 0 \rangle$ ).

*Spiking transition matrix*  $M_\Pi$  is a matrix comprised of  $a_{ij}$  elements where  $a_{ij}$  is given as

$$a_{ij} = \begin{cases} -c, & \text{if rule } r_i \text{ is in } \sigma_j \text{ and it is applied consuming } c \text{ spikes;} \\ p, & \text{if rule } r_i \text{ is in } \sigma_s \text{ (} s \neq j \text{ and } (s, j) \in \text{syn} \text{)} \\ & \text{and it is applied producing } p \text{ spikes in total;} \\ 0, & \text{if rule } r_i \text{ is in } \sigma_s \text{ (} s \neq j \text{ and } (s, j) \notin \text{syn} \text{)}. \end{cases}$$

For  $\Pi$ , the  $M_\Pi$  is as follows:

$$M_\Pi = \begin{pmatrix} -1 & 1 & 1 \\ -2 & 1 & 1 \\ 1 & -1 & 1 \\ 0 & 0 & -1 \\ 0 & 0 & -2 \end{pmatrix} \quad (1)$$

In such a scheme, rows represent rules and columns represent neurons.

Finally, the following equation provides the configuration vector at the  $(k+1)th$  step, given the configuration vector and spiking vector at the  $kth$  step, and  $M_\Pi$ :

$$C_{k+1} = C_k + S_k \cdot M_\Pi. \quad (2)$$

### 3. THE NVIDIA CUDA ARCHITECTURE

NVIDIA, a well known manufacturer of GPUs, released in 2006 the CUDA programming model and architecture [9]. Using extensions of the widely known C language, a programmer can write parallel code which will then execute in multiple threads within multiple thread blocks, each contained within a grid of (thread) blocks. These grids belong to a single device i.e. a single GPGPU. Each device/GPGPU has multiple cores, each capable of running its own grids. The program run in the CUDA model scales up or down, depending on the number of cores the programmer currently has in a device. This scaling is done in a manner that is abstracted from the user, and is efficiently as well. Automatic and efficient scaling is shown in Figure 2. Parallelized code will run faster with more cores than with fewer ones [8].

Figure 3 shows another important feature of the CUDA model: the host and the device parts. As mentioned earlier, device pertains to the GPGPU/s of the system, while the host pertains to the CPU/s. A function known as a kernel function, is a function called from the host but executed in the device.

A general model for creating a CUDA enabled program is shown in Listing 1.

#### Listing 1: General code flow for CUDA programming

```

1 //allocate memory on GPU e.g.
2 cudaMalloc( (void**)&dev_a, N * sizeof(int)
3
4 //populate arrays
5 . . .
6
7 //copy arrays from host to device e.g.
```

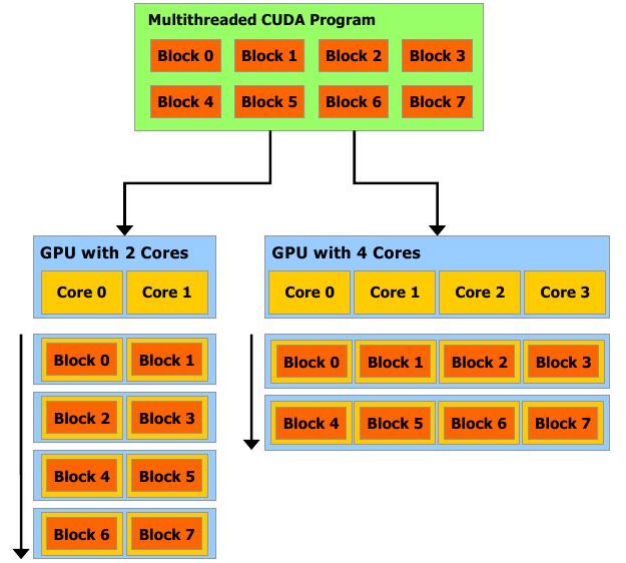


Figure 2: NVIDIA CUDA automatic scaling, hence more cores result to faster execution, from [8].

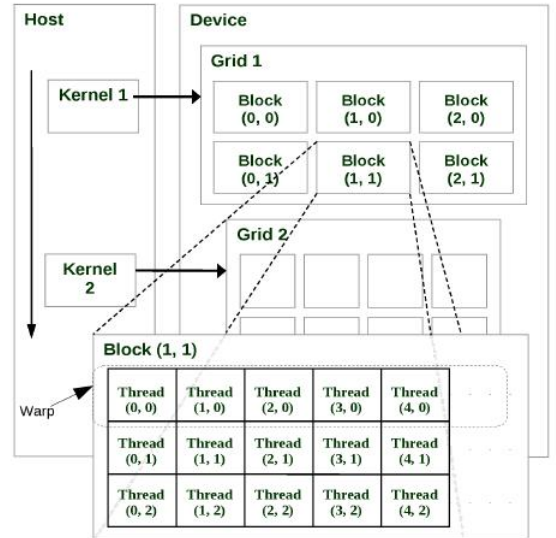


Figure 3: NVIDIA CUDA programming model showing the sequential execution of the *host* code alongside the parallel execution of the *kernel* function on the *device* side, from [6].

```

8 cudaMemcpy( dev_a, a, N * sizeof(int),
9 cudaMemcpyHostToDevice)
10
11 //call kernel (GPU) function e.g.
12 add<<<N, 1>>>( dev_a, dev_b, dev_c );
13
14 // copy arrays from device to host e.g.
15 cudaMemcpy( c, dev_c, N * sizeof( int),
16 cudaMemcpyDeviceToHost )
17
18 //display results
19
20 //free memory e.g.
21 cudaFree( dev_a );

```

Lines 2 and 19, implement CUDA versions of the standard C language functions e.g. the standard C function *malloc* has the CUDA C function counterpart being *cudaMalloc*, and the standard C function *free* has *cudaFree* as its CUDA C counterpart.

Lines 8 and 15 show a CUDA C specific function, namely *cudaMemcpy*, which, given an input of pointers ( from Listing 1 host code pointers are single letter variables such as *a* and *c*, while device code variable counterparts are prefixed by *dev\_* such as *dev\_a* and *dev\_c* ) and the size to copy ( as computed by the *sizeof* function ), moves data from host to device ( parameter *cudaMemcpyHostToDevice* ) or device to host ( parameter *cudaMemcpyDeviceToHost* ).

A kernel function call uses the double < and > operator, in this case the kernel function

*add << N, 1 >> (dev\_a, dev\_b, dev\_c).*

This function adds the values, per element (and each element is associated to 1 thread), of the variables *dev\_a* and *dev\_b* sent to the device, collected in variable *dev\_c* before being sent back to the host/CPU. The variable *N* in this case allows the programmer to specify *N* number of threads which will execute the *add* kernel function in parallel, with 1 specifying only one block for all *N* threads.

### 3.1 Design considerations given hardware and software setup

The kernel function i.e. code that is executed in parallel in the device, needs to have its results initially moved from the CPU/host to the device, and then back from the device to the host after computation. This movement of data back and forth should be minimized in order to obtain more efficient, in terms of time, execution. Implementing an equation such as 2, which involves multiplication and addition between vectors and a matrix, can be done in parallel with the previous considerations in mind. In this case,  $C_k$ ,  $S_k$ , and  $M_{\Pi}$  are loaded, manipulated, and pre-processed within the host code, before being sent to the kernel function which will perform computations on these function arguments in parallel. In order to represent  $C_k$ ,  $S_k$ , and  $M_{\Pi}$ , text files are created in order to house each input, whereby each element of the vector or matrix is entered in the file in order, from left to right, with a blank space in between as a delimiter. The matrix however is entered in row-major ( a linear array

of all the elements, rows first, then columns) order format i.e. for the matrix  $M_{\Pi}$  seen in 1, the row-major order version is simply

$$-1, 1, 1, -2, 1, 1, 1, -1, 1, 0, 0, -1, 0, 0, -2 \quad (3)$$

Row major ordering is a well-known ordering and representation of matrices for their linear as well as parallel manipulation in corresponding algorithms [7]. Once all computations are done for the  $(k+1)$ th configuration, the result of equation 2 are then collected and moved back from the device back to the host, where they can once again be operated on by the host/CPU. It is also important to note that these operations in the host/CPU provide logic and control of the data/inputs, while the device/GPU provides the arithmetic or computational 'muscle', the laborious task of working on multiple data or instructions at a given time, hence the current dichotomy of the CUDA programming model [6]. This division of labor is implemented in exemplified in Listing 1.

### 3.2 Matrix computations and CPU-GPGPU interactions

Once all 3 initial and necessary inputs are loaded, as is to be expected from equation 2, the device is first instructed to perform multiplication between the spiking vector  $S_k$  and the matrix  $M_{\Pi}$ . To further simplify computations at this point, the vectors are treated and automatically formatted by the host code to appear as single row matrices, since vectors can be considered as such. Multiplication is done per element (one element is in one thread of the device/GPGPU), and then the products are collected and summed to produce a single element of the resulting vector/single row matrix.

Once multiplication of the  $S_k$  and  $M_{\Pi}$  is done, the result is similarly added to the configuration vector  $C_k$ , once again element per element, with each element belonging to one thread, executed all at the same time as the others.

For this simulator, the host code consists almost entirely of the programming language Python, a well-known high- level, object oriented programming (OOP) language. The reason for using a high-level language such as Python is because the initial inputs, as well as succeeding ones resulting from exhaustively applying the rules and equation 2 require manipulation of the vector/matrix elements or values as strings to be concatenated, checked on (if they conform to the form (b-3) for example) by the host, as well as manipulated in ways which will be elaborated in the following sections along with the discussion of the algorithm for producing all possible and valid spiking vectors and configuration vectors given initial conditions. A language such as Python is well-suited for such a task, and can be byte-compiled like a C program for improved performance. The host code/Python part thus implements the logic and control as mentioned earlier, while in it, the device/GPU code which is written in C executes the parallel parts of the simulator.

## 4. DESIGN AND IMPLEMENTATION OF AN SNP SYSTEM SIMULATOR USING CUDA GPGPUS

The current SNP simulator, which is based on the type of SNP systems currently without time delays, is capable of implementing rules of the form (b-3) i.e. whenever the regular expression  $E$  is the same as the number of spikes consumed in that rule. Rules are entered in the same manner as the earlier mentioned vectors and matrix, as blank space delimited values (from one rule to the other, belonging to the same neuron) and \$ delimited ( from one neuron to the other). Thus for the SNP system  $\Pi$  shown earlier, the file  $r$  containing the blank space and \$ delimited values is as follows:

$$2 \ 2 \ \$ \ 1 \ \$ \ 1 \ 2 \quad (4)$$

That is, rule (1) from Figure 1 has the value 2 in the file  $r$  (though rule (1) isn't of the form (b-3) it nevertheless consumes a spike since its regular expression is of the same regular expression type as the rest of the rules of  $\Pi$ ). Another implementation consideration was the use of *lists* in Python, since unlike dictionaries or tuples, lists in Python are *mutable*, which is a direct requirement of the vector/matrix element manipulation to be performed later on (concatenation mostly). Hence a configuration vector  $C_k = \langle 2, 1, 1 \rangle$  is represented as  $[2, 1, 1]$  in Python. That is, at the  $k$ th configuration of the system, the number of spikes of neuron 1 are given by accessing the index (starting at zero) of the configuration vector list Python variable *confVec*, in this case if

$$\text{confVec} = [2, 1, 1] \quad (5)$$

then  $\text{confVec}[0] = 2$  are the number of spikes available at that time for neuron 1,  $\text{confVec}[1] = 1$  for neuron 2, and so on. The file  $r$ , which contains the ordered list of neurons and the rules that comprise each of them, is represented as a list of sub- lists in the Python/host code. For SNP  $\Pi$  we have the following:

$$r = [[2, 2], [1], [1, 2]] \quad (6)$$

Neuron 1's rules are given by accessing the sub-lists of  $r$  (again, starting at index zero) i.e. rule (1) is given by  $r[0][0] = 2$  and rule (4) is given by  $r[2][1] = 1$ .

## 4.1 Algorithms in the implementation of the SNP simulator

The general algorithm is as shown in Listing 2

Require: creation of files *confVec*, *M*, and *r* I. (host) Load inputs: configuration vector file ( *confVec* ), spiking transition matrix file (*M*), and rule criteria file ( *r* ). Note that *M* and *r* need only be loaded once since they are unchanging for a given SNP system.

II. (host) Determine if a rule/element in *r* is applicable based on the the spike value in

*configVec*, and then generate all valid + possible spiking vectors in a list of lists *spikVec* given *r* and *confVec*. III. ( device ) From part II, run the kernel function on *spikVec*, which contains all the valid + possible spiking vectors for the current *confVec* and *r*. This will generate further  $C_k$  and their corresponding  $S_k$ .

IV. Repeat steps I to IV until a zero configuration vector (vector with only zeros as elements) or further  $C_k$  produced are repetitions of a  $C_k$  produced at an earlier time.

## Listing 2: Overview of the algorithm for the SNP simulator

numbersepnumbersepnumbersep

## 4.2 Citations

Citations to articles [?, ?, ?, ?], conference proceedings [?] or books [?, ?] listed in the Bibliography section of your article will occur throughout the text of your article. You should use BibTeX to automatically produce this bibliography; you simply need to insert one of several citation commands with a key of the item cited in the proper location in the .tex file [?]. The key is a short reference you invent to uniquely identify each work; in this sample document, the key is the first author's surname and a word from the title. This identifying key is included with each item in the .bib file for your article.

The details of the construction of the .bib file are beyond the scope of this sample document, but more information can be found in the *Author's Guide*, and exhaustive details in the *LaTeX User's Guide*[?].

This article shows only the plainest form of the citation command, using \cite. This is what is stipulated in the SIGS style specifications. No other citation format is endorsed.

## 4.3 Tables

Because tables cannot be split across pages, the best placement for them is typically the top of the page nearest their initial cite. To ensure this proper "floating" placement of tables, use the environment **table** to enclose the table's contents and the table caption. The contents of the table itself must go in the **tabular** environment, to be aligned properly in rows and columns, with the desired horizontal and vertical rules. Again, detailed instructions on **tabular** material is found in the *LaTeX User's Guide*.

Immediately following this sentence is the point at which Table 1 is included in the input file; compare the placement of the table here with the table in the printed dvi output of this document.

To set a wider table, which takes up the whole width of the page's live area, use the environment **table\*** to enclose the table's contents and the table caption. As with a single-column table, this wide table will "float" to a location deemed more desirable. Immediately following this sentence is the point at which Table 2 is included in the input file; again, it is instructive to compare the placement of the table here with the table in the printed dvi output of this document.

## 4.4 Figures

Like tables, figures cannot be split across pages; the best placement for them is typically the top or the bottom of the page nearest their initial cite. To ensure this proper

Table 1: Some Typical Commands

Command	A Number	Comments
<code>\alignauthor</code>	100	Author alignment
<code>\numberofauthors</code>	200	Author enumeration
<code>\table</code>	300	For tables
<code>\table*</code>	400	For wider tables

Figure 4: A sample black and white graphic (.eps format).

Figure 5: A sample black and white graphic (.eps format) that has been resized with the epsfig command.

“floating” placement of figures, use the environment **figure** to enclose the figure and its caption.

This sample document contains examples of .eps and .ps files to be displayable with L<sup>A</sup>T<sub>E</sub>X. More details on each of these is found in the *Author’s Guide*.

As was the case with tables, you may want a figure that spans two columns. To do this, and still to ensure proper “floating” placement of tables, use the environment **figure\*** to enclose the figure and its caption.

Note that either .ps or .eps formats are used; use the `\epsfig` or `\psfig` commands as appropriate for the different file types.

## 4.5 Theorem-like Constructs

Other common constructs that may occur in your article are the forms for logical constructs like theorems, axioms, corollaries and proofs. There are two forms, one produced by the command `\newtheorem` and the other by the command `\newdef`; perhaps the clearest and easiest way to distinguish them is to compare the two in the output of this sample document:

This uses the **theorem** environment, created by the `\newtheorem` command:

THEOREM 1. Let  $f$  be continuous on  $[a, b]$ . If  $G$  is an antiderivative for  $f$  on  $[a, b]$ , then

$$\int_a^b f(t)dt = G(b) - G(a).$$

The other uses the **definition** environment, created by the `\newdef` command:

Definition 1. If  $z$  is irrational, then by  $e^z$  we mean the unique number which has logarithm  $z$ :

$$\log e^z = z$$

Two lists of constructs that use one of these forms is given in the *Author’s Guidelines*.

Figure 6: A sample black and white graphic (.ps format) that has been resized with the psfig command.

and don’t forget to end the environment with `figure*`, not `figure`!

There is one other similar construct environment, which is already set up for you; i.e. you must *not* use a `\newdef` command to create it: the **proof** environment. Here is an example of its use:

PROOF. Suppose on the contrary there exists a real number  $L$  such that

$$\lim_{x \rightarrow \infty} \frac{f(x)}{g(x)} = L.$$

Then

$$l = \lim_{x \rightarrow c} f(x) = \lim_{x \rightarrow c} \left[ g(x) \cdot \frac{f(x)}{g(x)} \right] = \lim_{x \rightarrow c} g(x) \cdot \lim_{x \rightarrow c} \frac{f(x)}{g(x)} = 0 \cdot L = 0,$$

which contradicts our assumption that  $l \neq 0$ .  $\square$

Complete rules about using these environments and using the two different creation commands are in the *Author’s Guide*; please consult it for more detailed instructions. If you need to use another construct, not listed therein, which you want to have the same formatting as the Theorem or the Definition[?] shown above, use the `\newtheorem` or the `\newdef` command, respectively, to create it.

## A Caveat for the T<sub>E</sub>X Expert

Because you have just been given permission to use the `\newdef` command to create a new form, you might think you can use T<sub>E</sub>X’s `\def` to create a new command: *Please refrain from doing this!* Remember that your L<sup>A</sup>T<sub>E</sub>X source code is primarily intended to create camera-ready copy, but may be converted to other forms – e.g. HTML. If you inadvertently omit some or all of the `\defs` recompilation will be, to say the least, problematic.

## 5. CONCLUSIONS AND FUTURE WORK

Using a highly parallel computing device such as a GPGPU, particularly NVIDIA CUDA, an SNP system simulator was successfully designed and implemented. The use of a high level programming language such as Python for host tasks, mainly for logic and string representation and manipulation of values (vector/matrix elements) provided the necessary expressivity to implement the algorithms created to produce and exhaust all possible and valid configuration and spiking vectors. For the device tasks, CUDA C allowed the manipulation of the NVIDIA CUDA enabled GPGPU which took

**Figure 7: A sample black and white graphic (.eps format) that needs to span two columns of text.**

care of repetitive and highly parallel computations (addition and multiplication essentially).

Future versions of the SNP system simulator will focus on several improvements. These improvements include the use of an algorithm for matrix computations without requiring the input matrix to be turned into a square matrix (this is currently handled by the simulator by padding zeros to an otherwise non-square matrix input). Another improvement would be the simulation of systems not of the form  $b-3$ ). Byte-compiling the Python/host part of the code to improve performance as well as metrics to further enhance and measure execution time are desirable as well. Finally, deeper understanding of the CUDA architecture, such as inter-thread/block communication, for extremely large systems with equally large matrices, is required. These improvements as well as the current version of the simulator should also be run in a machine with higher versions of GPG-Us running NVIDIA CUDA.

## 6. ACKNOWLEDGMENTS

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