# Accelerator-Oriented Algorithm Transformation for Temporal Data Mining 

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#### Abstract

Temporal data mining algorithms are becoming increasingly important in many application domains including computational neuroscience, especially the analysis of spike train data. While application scientists have been able to readily gather multi-neuronal datasets, analysis capabilities have lagged behind, due to both lack of powerful algorithms and inaccessibility to powerful hardware platforms. The advent of GPU architectures such as Nvidia's GTX 280 offers a costeffective option to bring these capabilities to the neuroscientist's desktop. Rather than port existing algorithms onto this architecture, we advocate the need for algorithm transformation, i.e., rethinking the design of the algorithm in a way that need not necessarily mirror its serial implementation strictly. We present a novel implementation of a frequent episode discovery algorithm by revisiting "in-the-large" issues such as problem decomposition as well as "in-the-small" issues such as data layouts and memory access patterns. This is non-trivial because frequent episode discovery does not lend itself to GPU-friendly data-parallel mapping strategies. Applications to many datasets and comparisons to CPU as well as prior GPU implementations showcase the advantages of our approach.


Keywords-GPGPU; Temporal data mining; Frequent episodes; Spike train analysis; Computational neuroscience; CUDA

## I. Introduction

Discovering frequently repeating patterns in event sequences is an important data mining problem that finds application in domains such as industrial plants/assembly lines, medical diagnostics, and computational neuroscience. Algorithms such as frequent episode discovery [1], [2], in particular, are adept at discovering patterns in neuronal spike train data using multi-electrode arrays (MEAs; shown in Fig. 1). They bring us one step closer to reverse-engineering the temporal connectivity map of neuronal circuits and yield insights into the network level activity of brain tissue. Until recently, most analysis was limited to single cell activity due to the low computational capacity of single-core computers and lack of efficient data mining algorithms. While clever data mining algorithms have made it feasible to overcome the combinatorial explosion problem of discovering multiway interactions, they are still limited by the memory and processing power constraints of the single CPU.

In recent years, the peak-performance of a GPU has exceeded that of the CPU by several orders of magnitude. Intel's latest quad-core processors have a theoretical peak of 51.20 GFLOPS. Nvidia's latest single GPU card, the GeForce GTX 285, has a theoretical peak of 1062.72


Figure 1. Illustration of a Micro electrode array (MEA) used to record spiking activity of neurons in tissue cultures and the spike train recorded by the MEA.

GFLOPS. Application speedups up to 431x have been reported [3]. The demand for graphics applications like gaming have made GPUs widely available and inexpensive. In this paper we explore how this massively parallel computing platform can be used effectively to solve the challenges posed by temporal data mining. Although we focus on a specific algorithm, the issues encountered here are symptomatic of many temporal mining algorithms that must use state machines to monitor and process event streams. Hence the lessons learned from this effort will likely seed similar research efforts.

One major challenge in utilizing the GPU lies in transforming algorithms to operate efficiently on the GPU. Another challenge is understanding how to use the GPU architecture to achieve maximum performance. Data-parallel algorithms require relatively less work to map computation onto the GPU architecture. For algorithms with complex data dependencies, such as dynamic programming, it is difficult to achieve a large speedup. The nature of the temporal data mining problem addressed in this work limits the performance gain of standard hardware-oriented optimizations applied to direct ports of existing algorithms (due to their sequential dependencies implicit in processing event streams).

In this paper, we adopt the concept of AcceleratorOriented Algorithm Transformation and introduce a new algorithm for counting occurrences of frequent episodes with temporal constraints (a key analysis task for event stream analysis) on the GPU. Here an episode is a sequential dependency of the form 'event A followed by event B followed by event C...' where there could be don't care or "junk" events interspersed between the pattern events. (The frequency of such episodes is defined as the maximum number of non-overlapped occurrences of the episodes in the event stream.)

The traditional approach to parallelizing existing algorithms is to start with the existing sequential algorithm, identify the data dependency patterns, and restructure the original algorithm to achieve maximum data parallelism. Another way is to look at the problem being solved and formulate a decomposition using known parallel primitives. We follow the later approach and design a new algorithm by decomposing the original problem into two sub-problems: finding overlapped episode occurrences and resolving overlaps to obtain non-overlapped counts. We introduce a parallel local tracking algorithm to solve the first sub-problem which is computationally more demanding. On the other hand, the second sub-problem contributes only a very small overhead to the overall computation and is hence solved sequentially. The re-designed algorithm exhibits a higher degree of parallelism resulting in a performance gain over the sequential algorithm implemented on a single-core CPU and a previously optimized GPU implementation (MapConcat) [4] which achieves parallelism by mining segments of the data sequence.

## II. Background

## A. GPGPU Architecture

To understand the algorithmic details behind our approach, we provide a brief overview of GPGPU and its architecture.

The initial purpose of specialized GPUs was to accelerate the display of 2D and 3D graphics, much in the same way that the FPU focused on accelerating floating-point instructions. However, the rapid technological advances of the GPU, coupled with extraordinary speed-ups of application "toy" benchmarks on the specialized GPUs, led GPU vendors to transform the GPU from a specialized processor to a general-purpose graphics processing unit (GPGPU), such as the NVIDIA GTX 280. To lessen the steep learning curve, GPU vendors also introduced programming environments, such as the Compute Unified Device Architecture (CUDA).


Figure 2. The NVIDIA GTX280 GPU architecture.
Processing Elements: The basic execution unit on the GTX 280 is a scalar processing core, of which 8 together form a multiprocessor. While the number of multiprocessors and processor clock frequency depends on the make and model
of the GPU, every multiprocessor in CUDA executes in SIMT (Single Instruction, Multiple Thread) fashion, which is similar in nature to SIMD (Single Instruction, Multiple Data) execution. Specifically, a group of 32 threads form a warp and are scheduled to execute concurrently. However, when codepaths within a warp diverge, the execution of all threads in a warp becomes serialized. This implies that optimal performance is attained when all 32 threads do not branch down different codepaths.
Memory Hierarchy: The GTX 280 contains multiple forms of memory. The read-write global memory and read-only texture memory is located off-chip on the graphics card and provides the main source of storage for the GPU, as shown in Figure 2. Each multiprocessor on the GPU contains fast on-chip memory, which includes cache memory and shared memory. The texture cache is read-only memory providing fast access to immutable data. Shared memory, on the other hand, is user-controlled read-write space to provide each core with fast access to the shared address space within a multiprocessor.
Parallelism Abstractions: At the highest level, the CUDA programming model requires the programmer to offload functionality to the GPGPU as a compute kernel. This kernel is evaluated as a set of thread blocks logically arranged in a grid to facilitate organization. In turn, each block contains a set of threads, which will be executed on the same multiprocessor, with the threads scheduled in warps, as mentioned above.

## B. Data Mining using GPGPUs

Many researchers have harnessed the capabilities of GPGPUs for data mining. The key to porting a data mining algorithm onto a GPGPU is to, in one sense, "dumb it down"; i.e., conditionals, program branches, and complex decision constraints are not easily parallelizable on a GPGPU and algorithms using these constraints will require significant reworking to fit this architecture (temporal episode mining unfortunately falls in this category). There are many emerging publications in this area but due to space restrictions, we survey only a few here. The SIGKDD tutorial by Guha et al. [5] provides a gentle introduction to the aspects of data mining on GPGPUs through problems like k-means clustering, reverse nearest neighbor(RNN), discrete wavelet transform(DWT), sorting networks, etc. In [6], a bitmap technique is proposed to support counting and is used to design GPGPU variants of Apriori and k-means clustering. This work also proposes co-processing for itemset mining where the complicated tie data structure is kept and updated in the main memory of CPU and only the itemset counting is executed in parallel on GPU. A sorting algorithm on GPGPUs with applications to frequency counting and histogram construction is discussed in [7] which essentially recreates sorting networks on the GPU.

## C. Temporal Data Mining

In event sequences, the notion of frequent episodes is used to express patterns of the form $\mathrm{A} \rightarrow \mathrm{B} \rightarrow \mathrm{C}$, i.e., event A is followed (not necessarily consecutively) by B and B is, similarly, followed by C. It is important to constrain the mining by imposing minimum and maximum delays between consecutive symbols in an episode, e.g., to look for episodes of the form $(A \xrightarrow{(2,5]} B \xrightarrow{(0,6]} C)$. This specifies that event A is followed by B within two to five milliseconds, and C follows $B$ within at most six milliseconds. In either unconstrained or constrained episode mining, the occurrences of an episode allow 'junk' or don't care events, of arbitrary length, between the event symbols of the episode. This is what makes these patterns very useful and comprehensible. Many frequency measures [1], [8] for episodes have been defined that obey anti-monotonicity and hence search for such episodes can be structured level-wise, ala Apriori. The first measure to be proposed was the window based frequency measure [8]. Later the notion of non-overlapped occurrences count was shown to have properties that enable fast sequential counting algorithms [1].

Definition 1: An Event Stream is denoted by a sequence of events $\left\langle\left(E_{1}, t_{1}\right),\left(E_{2}, t_{2}\right), \ldots\left(E_{n}, t_{n}\right)\right\rangle$, where $n$ is the total number of events. Each event $\left(E_{i}, t_{i}\right)$ is characterized by an event type $E_{i}$ and a time of occurrence $t_{i}$. The sequence is ordered by time i.e. $t_{i} \leq t_{i+1} \forall i=1, \ldots, n-1$ and $E_{i}$ 's are drawn from a finite set $\xi$.

In neuroscience, a spike train is a series of discrete action potentials from several neurons generated spontaneously or as a response to some external stimulus. This data neatly fits into the frequent episodes framework of analyzing event streams.

Definition 2: An (serial) episode $\alpha$ is an ordered tuple of event types $V_{\alpha} \subseteq \xi$.
For example $(A \rightarrow B \rightarrow C)$ is a 3-node serial episode, and it captures the pattern that neuron A fires, followed by neurons B , and C in that order, but not necessarily consecutive.

Frequency of episodes: A frequent episode is one whose frequency exceeds a user specified threshold. The notion of frequency of an episode is intended to capture the repeating nature of an episode in an event sequence. In this work we shall focus on the measure of frequency defined as the size of the largest set of non-overlapped occurrences of an episode [1].

Temporal constraints: Episodes can be further specialized by specifying constraints on the timing of the events in episode occurrences [2]. Placing inter-event time constraints giving rise to episodes of the form:

$$
\left(A \xrightarrow{\left(t_{l o w}^{1}, t_{h i g h}^{1}\right]} B \xrightarrow{\left(t_{l o w}^{2}, t_{h i g h}^{2}\right]} C\right)
$$

That is, in an occurrence of episode $A \rightarrow B \rightarrow C$ let $t_{A}$, $t_{B}$, and $t_{C}$ denote the time of occurrence of corresponding
event types. A valid occurrence of the serial episode satisfies

$$
t_{l o w}^{1}<\left(t_{B}-t_{A}\right) \leq t_{\text {high }}^{1}, t_{l o w}^{2}<\left(t_{C}-t_{B}\right) \leq t_{h i g h}^{2}
$$

(In general, an $N$-node serial episode is associated with $N-$ 1 inter-event constraints.)

Level-wise discovery procedure for frequent episodes finds the complete set of episodes with frequency or count greater than a user-defined threshold. At each level $(N)$ size candidates are generated from $(N-1)$-size frequent episodes and their count is determined by making a pass over the event sequence. Only those candidates whose count is greater than the threshold are retained. The event sequences typically runs very long and hence the counting step is computationally most expensive. For initial passes of the algorithm where we have several short episodes to count the counting task can be embarrassingly parallel by making each thread of execution count occurrences of only one episode. But in later stages there are relatively fewer but longer episodes leading to severe under-utilization. We focus our work here on the later situation and attempt to solve this counting problem.

## III. Algorithm

## A. Episode Counting Algorithm

The algorithm presented in [2] is based on finite state machines. This sequential algorithm for mining frequent episodes with inter-event constraints works by maintaining a data-structure shown in Figure 3 as the read head moves down the event sequence. In this example we are counting occurrences of episode $A \xrightarrow{(5,10]} B \xrightarrow{(10,15]} C$.


Figure 3. Illustration of the algorithm for counting non-overlapped occurrences of $A \xrightarrow{(5,10]} B \xrightarrow{(10,15]} C$. This proceeds from left-to-right of the event steam keeping track of sufficient information required to obtain the correct non-overlapped count under the given inter-event time constraints.

The general approach, on finding an event that belongs to the episode, is to look up the list of occurrences of the previous symbol (or event-type). If there exists an event of the previous event type which together with the new event satisfies the inter-event constraint for the pair of eventtypes, then the new event is added to the data-structure under its corresponding symbol. An occurrence is said to be complete when we can add an event for the last symbol
in the episode. Then the count is incremented and the datastructure cleared. For example when $(B, 18)$ arrives, it is found that the pair $(A, 10) \in \operatorname{list}[A]$ and $(B, 18)$ satisfy the inter-event constraint $(5,10]$. And therefore the time for $(B, 18)$ is recoded in list $[B]$. On arrival of $(C, 20)$ we are able to complete one occurrence of the episode.

## B. MapConcat

The initial idea for parallelization is to map one thread per episode. However, when the number of episodes is smaller than the number of cores, a one-thread per-episode model is prone to under utilization leading to higher execution times. Hence our first attempt to achieve a higher level of parallelism within the counting of a single episode was to segment the input event stream into several of sub-streams and use one thread block to count one episode [4].


Figure 4. Illustration of a Map step. Multiple state machines are needed to track occurrences of $N$-size episode $\alpha$ in the $p^{t h}$-Segment of the data. Each state machine starts at a different offset into the previous segment and continues over into the next segment to complete the last occurrence. The end-time of the first $a$ and last occurrence $b$ seen by a state machine are recorded beside the count for the reduce/concat step.

When we divide the input stream into segments, there are chances that some occurrences of an episode span across the boundaries of consecutive segments. It turns out that in order to obtain the correct count we are required to run multiple state machines within the same data segment anticipating all possible end states of the state machine in the previous segment. The final count is obtained by a reduce step where state machines for consecutive segments with matching start and end states are concatenated. The counting step (or the Map-step) is illustrated in Fig. 4 and the reduce step (or Concat-step) is shown in Fig. 5

## IV. Redesigned Algorithm

The original state-machine based algorithm has a lot of data dependency. It is difficult to increase the degree of parallelism by optimizing this algorithm. In order to transform the algorithm to map well onto the GPGPU architecture, we must revisit the problem and formulate a decomposition using known parallel primitives.

## A. Problem decomposition

Referring to the definition of non-overlapped frequency of an episode, it is the size of the largest set of non-overlapped


Figure 5. Illustration of a Concatenate step. As noted in Fig 4 the first and last occurrence of each state-machine is used to stitch together the total count for an episode. The blue arrow in this figure indicates that the last occurrence seen by the state-machine on the left matches with the first occurrence seen by the state-machine on the right. These state-machines can then be merged together into the next level.
occurrences. Based on this definition we design a more data parallel solution. In our new approach we find a super-set of the non-overlapped occurrences which could potentially be overlapping. Each occurrence of an episode has fixed start and end times. If each episode occurrence in this super set can be viewed as a task with start and end time, then the problem of finding the largest set of non-overlapped occurrences can be easily transformed into a task scheduling problem, where the goal is to maximize number of nonconflicting tasks on a single shared resource. It is known that a greedy $O(n)$ algorithm can solve this problem optimally. The original problem now decomposes into the following two sub-problems:

- Subproblem-1: Finding a super-set of non-overlapped occurrences of an episode.
- Subproblem-2: Finding the size of the largest set of non-overlapped occurrences from the above set of occurrences.
The first sub-problem can be solved with high parallelism as will be shown in following sections. The second subproblem is same as the task or interval scheduling problem where tasks have fixed times. A fast greedy algorithm is known to solve the problem optimally.

We first pre-process the entire event stream noting the positions of events of each event-type. Then for a given episode, beginning with the list of occurrences of the start event-type in the episode, we find occurrences satisfying the temporal constraints in parallel. Finally we collect and remove overlapped occurrences in one pass. The greedy algorithm for removing overlaps requires the occurrences to be sorted by end time and the algorithm proceeds as shown in Algorithm 1. In this algorithm, for every set of consecutive occurrences, if the start time is after the end time of the last selected occurrence then we select this occurrence, otherwise we skip it and go to the next occurrence.

## B. Finding occurrences in parallel

Here we explore different approaches of solving the first sub-problem as discussed in the previous section. The aim

```
Algorithm 1 Obtaining the largest set of non-overlapped
occurrences
Input: List \(\mathcal{C}\) of occurrences with start and end times \(\left(s_{i}, e_{i}\right)\)
    sorted by end time, \(e_{i}\).
Output: Size of the largest set of non-overlapped occur-
    rences
    Initialize count \(=0\)
    preve \(_{e}=0\)
    for all \(\left(s_{i}, e_{i}\right) \in \mathcal{C}\) do
        if prev \(_{e}<s_{i}\) then
            prev \(_{e}=e_{i} ;\) count \(=\) count +1
    return count
```

here is to find a super-set of non-overlapped occurrences in parallel. The basic idea is to start with all events of the first event-type in parallel for a given episode and find occurrences of the episode starting at each of these events. There can be several different ways in which this can be done. We shall present two approaches that gave us the most performance improvement. We shall use the episode $A \xrightarrow{(5-10]} B \xrightarrow{\text { (5-10] }} C$ as our running example and explain each of the counting strategies using this example. This example episode specifies event occurrences where an event A is to be followed by an event $B$ within $5-10 \mathrm{~ms}$ and event $B$ is to be followed by an event $C$ within $5-10 \mathrm{~ms}$ delay. Note again that the delays have both a lower and a upper bound.

## C. Parallel Local Tracking

In the pre-processing step, we have noted the locations of each of the event-types in the data. In the counting step, we launch as many threads as there are events of the start type. In our running example this is all events of type $A$. Each thread searches the event streams starting at one of these events of type $A$ and looks for an event of type $B$ that satisfies the inter-event time constraint $(5-10]$ i.e., $5<t_{B_{j}}-t_{A_{i}} \leq 10$ where $i, j$ are the indices of the events of type $A$ and $B$. One thread can find multiple $B$ 's for the same $A$. These are recorded in a preallocated array assigned to each thread. Once all the events of type $B$ (with an $A$ before them) have been collected by the threads (in parallel), we need to compact these newly found events into a contiguous array/list. This is necessary as in the next kernel launch we will find all the events of type $C$ that satisfy the inter-event constraints with this set of $B$ 's. This is illustrated in Figure 6

Algorithm 2 presents the over-all work done in each kernel launch. In order to obtain the complete set of occurrences of an episode, we need to launch the kernel $N-1$ times where $N$ is the size of an episode. The list of events found in the $i^{t h}$ iteration is passed as input to the next iteration. Some amount of book-keeping is also done to keep track of the start and end times of an occurrence. After this phase of parallel local tracking is completed the non-overlapped


Figure 6. Illustration of Parallel local tracking algorithm (See Algorithm 2], showing 2 iterations for the episode $A \rightarrow B \rightarrow C$ with implicit inter-event constraints. Note that each thread can find multiple next-events. Further, a thread stops scanning the event sequence when event-times go past the upper bound of the inter-event constraint.

```
Algorithm 2 Kernel for Parallel Local Tracking
Input: Iteration number \(i\), Episode \(\alpha\), Event type \(\alpha[i]\), Index
    list \(I_{\alpha[i]}\), Data sequence \(S\).
Output: Indices of events of type \(\alpha[i+1]\).
    for all Threads which have distinct identifiers tid do
        Scan \(S\) starting at event \(I_{\alpha[i]}[t i d]\) for event-type \(\alpha[i+1]\)
        satisfying inter-event constraint \(\left(t_{\text {low }}^{(i)}, t_{\text {high }}^{(i)}\right]\).
        Record all such events of type \(\alpha[i+1]\).
    Compact all found events into the list \(I_{\alpha[i+1]}\).
    return \(I_{\alpha[i+1]}\)
```

count is obtained using Algorithm 1. The compaction step in Algorithm 2 presents a challenge as it requires concurrent updates into a global array.

## D. Lock-based compaction

Nvidia graphics cards with CUDA compute capability 1.3 support atomic operations on shared and global memory. Here we use atomic operations to perform compaction of the output array into the global memory. After the counting step each thread has a list of next-events. Subsequently each thread adds the size of its next-events list to the block-level counter using an atomic add operation and in return obtains a local offset (which is the previous value of the block-level counter). After all threads in a block have updated the blocklevel counter, the first thread in the block updates the globalcounter by adding the value of the block-level counter to it and, as before, obtains the offset into global memory. Now all threads in the block can collaboratively write into the correct position in the global memory (resulting in overall compaction). A schematic for this operation is shown for 2-blocks in Figure 7. In the results section, we refer to this method as AtomicCompact.

Since there is no guarantee for the order of atomic operations, this procedure requires sorting. The complete occurrences need to be sorted by end time for Algorithm 1 to produce the correct result.


Figure 7. Illustration of output compaction using AtomicAdd operations. Note that we use atomic operations at both block and global level. These operations return the correct offset into global memory for each thread to write its next-event list into.

## E. Lock-free compaction

Prefix-scan is known to be a general-purpose data-parallel primitive that is a useful building block for algorithms in a broad range of applications. Given a vector of data elements $\left[x_{0}, x_{1}, x_{2}, \ldots\right]$, an associative binary function $\oplus$ and an identity element $i$, exclusive prefix-scan returns [ $\left.i, x_{0}, x_{0} \oplus x_{1}, x_{0} \oplus x_{1} \oplus x_{2}, \ldots\right]$. Although the problem is seemingly sequential the first parallel prefix-scan algorithm was proposed in 1962 [ 9 ]. With recently increasing interest is GPGPU, several implementations of scan have been proposed for GPU, the most recent being [10], [11]. This later implementation is available as the CUDPP: CUDA Data Parallel Primitives Library and forms part of the CUDA SDK distribution.

Our lock-free compaction is based on prefix-sum and we reuse the implementation from CUDPP library. Since the scan based operation guarantees ordering we modify our counting procedure to count occurrences backwards starting from the last event. This results in the final set of occurrences to be automatically ordered by end-time and therefore completely eliminates the need for sorting (as required by atomic operations based approach).


Figure 8. Illustration of output compaction using Scan primitive. Each iteration is broken into 3 kernel calls: Counting the number of next events, Using scan to compute offset into global memory, finally launching countprocedure again but this time allowing write operations to the global memory

The CUDPP library provides a compact function which takes an array $d_{i n}$, an array of $1 / 0$ flags and returns a compacted array $d_{\text {out }}$ of corresponding only the "valid"
values from $d_{i n}$ (it internally uses cudppScan). In order to use this, our counting kernel is now split into three kernel calls. Each thread is allocated a fixed portion of a larger array in global memory for its next-events list. In the first kernel, each thread finds its events and fills up its nextevents list in global memory. The cudppCompact function (implemented as two GPU kernel calls) compacts the large array to obtain the global list of next-events. A difficulty of this approach is that the array on which cudppCompact operates is very large resulting in a scattered memory access pattern. We refer to this method as CudppCompact.

In order to improve performance, we adopt a counterintuitive approach. We divide the counting process into three parts. First each thread looks up the event sequence for suitable next-events but instead of writing the found events anywhere, it merely counts and writes the count to global memory. Then an exclusive scan is performed for the countarray which gives the offset into the global memory where each thread can write its next-events list. The actual writing is done as the third step. Although each thread looks up the event sequence twice (first to count, and second to write) we show that we nevertheless obtain better performance. This entire procedure is illustrated in Figure 8 . We refer to this method of compaction as CountScanWrite in the ensuing results section.

## V. Results

The hardware used for obtaining the performance results are given in Table II

Table I
HARDWARE USED FOR PERFORMANCE ANALYSIS

| GPU | Nvidia GTX 280 |
| :--- | :---: |
| Memory (MB) | 1024 |
| Memory Bandwidth (GBps) | 141.7 |
| Multiprocessors, Cores | 30,240 |
| Processor Clock (GHz) | 1.3 |
| CPU | Intel Core 2 Quad Q8200 |
| Processors | 4 |
| Processor Clock (GHz) | 2.33 |
| Memory (MB) | 4096 |

## A. Test datasets and algorithm implementations

The datasets used here are generated from the nonhomogeneous Poisson process model for inter-connected neurons described in [2]. This simulation model generates fairly realistic spike train data. For the datasets in this paper a networks of 64 artificial neurons was used. The random firing rate of each neuron was set at 20 spikes/sec to generate sufficient noise in the data. Four 9-node episodes were embedded into the network by suitably increasing the connection strengths for pairs of neurons. Spike train data was generated by running the simulation model for different durations of time. Table $\Pi$ gives the duration and number of events in each dataset.

Table II
DETAILS OF THE DATASETS USED

| Data <br> -Set | Length <br> (in sec) | \# Events |
| :---: | :---: | :---: |
| 1 | 4000 | $12,840,684$ |
| 2 | 2000 | $6,422,449$ |
| 3 | 1000 | $3,277,130$ |
| 4 | 500 | $1,636,463$ |

In Section V-B, we compare the performance of our new algorithm to MapConcat and a CPU implementation of the original algorithm described in Section II-C. In order to analyze the lock-based and lock-free compaction strategies, we present the performance of a lock-based method, AtomicCompact, and two lock-free methods, CudppCompact and CountScanWrite, as shown in Section V-C

## B. Comparisons of performance



Figure 9. Performance of MapConcat compared with the CPU and best GPU implementation, counting 30 episodes in Datasets 1-8.

MapConcat is clearly a poor performer compared to the CPU, with up to a 4 x slowdown. Compared to our best GPU method, MapConcat is up to 11 x slower. This is due to the overhead induced by the merge step of MapConcat. Although at the multiprocessor level each episode is counted in parallel, the logic required to obtain the correct count is complex.

We run the CUDA Visual Profiler on MapConcat and one of our redesigned algorithms, CountScanWrite. Dataset 2 was used for profiling each implementation. Due to its complexity, MapConcat exhibited poor features such as large amounts of divergent branching and a large total number of instructions executed, as shown in Table III Comparatively, the CountScanWrite implementation only exhibits divergent branching.

Table III
CUDA Visual Profiler Results

|  | MapConcat | CountScanWrite |
| :--- | :---: | :---: |
| Instructions | $93,974,100$ | $8,939,786$ |
| Branching | $27,883,000$ | $2,154,806$ |
| Divergent Branching | $1,301,840$ | 518,521 |



Figure 10. Performance comparison of the CPU and best GPU implementation, counting a single episode in Datasets 1 through 8.

In terms of the performance of our best GPU method, we achieve a $6 x$ speedup over the CPU implementation on the largest dataset, as shown in Figure 10 .

## C. Analysis of the new algorithm



Figure 11. Performance of algorithms with varying episode length in Dataset 1.

Figure 11 contains the timing information of three compaction methods of our redesigned GPU algorithm with varying episode length. Compaction using CUDPP is the slowest of the GPU implementations, due to its method of compaction. It requires each data element to be either in or out of the final compaction, and does not allow for compaction of groups of elements. For small episode lengths, the CountScanWrite approach is best because sorting can be completely avoided. However, at longer episode lengths, compaction using lock-based operators shows the best performance. This method of compaction avoids the need to perform a scan and a write at each iteration, at the cost of sorting the elements at the end. The execution time of the AtomicCompact is nearly unaffected by episode length, which seems counter-intuitive because each level requires a kernel launch. However, each iteration also decreases the total number of episodes to sort and schedule at the end of the algorithm. Therefore, the cost of extra kernel invocations is offset by the final number of potential episodes to sort and schedule.


Figure 12. Performance of algorithms with varying episode frequency in Dataset 1.

We find that counting time is related to episode frequency as shown in Figure 12. There is a linear trend, with episodes of higher frequency require more counting time. The lock-free compaction methods follow an expected trend of slowly increasing running time because there are more potential episodes to track. The method that exhibits an odd trend is the lock-based compaction, AtomicCompact. As the frequency of the episode increases, there are more potential episodes to sort and schedule. The running time of the method becomes dominated by the sorting time as the episode frequency increases.

Another feature of Figure 12 that requires explanation is the bump where the episode frequecy is slightly greater than 80,000 . This is because it is not the final non-overlapped count that affects the running time, it is the total number of overlapped episodes found before the scheduling algorithm is appiled to remove overlaps. The x -axis is displaying nonoverlapped episode frequency, where the run-time is actually affected more by the overlapped episode frequency.

We used the CUDA Visual Profiler on the other GPU methods. They had similar profiler results as the CountScanWrite method. The reason is that the only bad behavior exhibited by the method is divergent branching, which comes from the tracking step. This tracking step is common to all of the GPU method of the redesigned algorithm.

## VI. Conclusion

Just as algorithms for secondary storage are quite distinct from algorithms for main memory (e.g., two-phase merge sort is preferable over quicksort), we have shown similarly that approaches for temporal data mining on a GPU must adopt fundamentally different strategies than that on a CPU. Through this work, we aim to have conveyed that even with an application such as frequent episode mining that is uneqivocally 'sequential' in nature, it is possible to obtain reasonable speedup by an order of magnitude using careful redesign and algorithm-oriented transformation. A key lesson from our efforts is the importance of investigating both "in-the-large" and "in-the-small" issues. We have demonstrated the effectiveness of our approach to handling
large scale event stream datasets modeled by inhomogeneous Poisson processes.

## VII. Future Work

Similar to our motivations stemming from computational neuroscience, there are a large class of data mining tasks in bioinformatics, linguistics, and event stream analysis that require analysis of sequential data. Our work opens up the interesting issue of the extent to which finite state-machine based algorithms for these tasks can be accelerated using GPU platforms. Are there fundamental limitations to porting such algorithms on GPUs? We believe there are and hope to develop a theoretical framework to investigate GPUtransformation issues for these algorithms. Second, we aim to study the development of streaming versions of these algorithms where approximate results are acceptable but near real-time responsiveness is important.

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