

ARTICLE TYPE

Fast LRT Implementation on Parallel Computer Architectures

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Summary

The Likelihood Ratio Test (LRT) is a method for identifying hotspots or anomalous rectangular regions for an spatial data grid that draws from an arbitrary distribution. LRT has wide-spread applications from social-media applications powered by mobile phones to medical image processing applications. Unfortunately, a naïve implementation of LRT exhibits a worst-case time complexity of $\mathcal{O}(n^4)$ for a two-dimensional grid. For larger grids with a grid size of more than 64×64 cells, a naïve implementation of LRT becomes intractable (1) and it takes around several days to execute Grid(64,64) on a single-core CPU.

In this paper, we propose a parallel processing scheme for LRT computations for one parameter exponential (1EXP) distribution for multi-dimensional spatial data grids. The parallel processing scheme transforms LRT computations to a completely balanced embarrassingly parallel problem utilising a novel range mapping scheme. The new range mapping scheme relies on a dynamic pre-computation algorithm, and exhibits a worst-case complexity of $\mathcal{O}(n^2)$ for two dimensional spatial data-grids improving the complexity class by an order of two. Experiments were conducted for various computer architectures and platforms including shared-memory multi-core machines, multi-GPGPU computers, and the EC2 cloud computer. Extensive experiments are provided to demonstrate the utility of this approach and extensive performance analysis is given. In the experiments we could achieve speedups up to 20,000 times using our parallel processing scheme on Grid (1000, 1000).

KEYWORDS:

Spatial outlier, Likelihood Ratio Test, one parameter exponential distribution, Inclusive/Exclusive Principle, GPGPUs, Multi-core, EC2 Cloud Cluster

1 | INTRODUCTION

With the widespread availability of GPS-equipped smartphones and mobile sensors, there has been an urgent need to perform large scale spatial data analysis. For example, by carrying out a geographic projection of Twitter feeds, researchers are able to narrow down “hotspot” regions where a particular type of activity is attracting a disproportionate amount of attention. In neuroscience, high resolution MRIs facilitate the precise detection and localisation of regions of the brain which may indicate mental disorder. The statistical method of choice for identifying hotspots or anomalous regions is the Likelihood Ratio Test (LRT) statistic. Informally, the LRT of a spatial region compares the likelihood of the given spatial region with its complement, and hence can be used to identify hotspot regions. In (2), it was shown that the LRT value follows a χ^2 distribution, independent of the distribution of the underlying data.

For a $n \times n$ spatial grid, the worst-case execution time for identifying the most anomalous region is $\mathcal{O}(n^4)$. The reason for this high complexity class is that we have $\mathcal{O}(n^2)$ regions in the grid and computing the value of a single region R in the grid consumes $\mathcal{O}(n^2)$. As noted by Wu et al.(1), a naive implementation of LRT for a moderately sized 64×64 spatial grid may take nearly six hundred days¹. Wu et al.(1) proposed a method which reduces the computation time to eleven days. However as noted previously in (3), this approach will not scale for larger data sets and the biggest spatial grid reported in(1) was 64×64 .

The nature of LRT permits the independent computation of regions. This facilitates parallelization to some degree. However, the LRT computation of a region R involves the irregularly shaped computation of \bar{R} (4). In a parallel environment, this can drastically reduce its computation performance(5, 6, 7). To reduce the overall enumeration cost and address the irregular computation, Pang et al. (4) presented a unified parallel approach for generalized LRT computation in spatial data grids on a GPGPU environment. The whole grid is partitioned into overlapped blocks and LRT computation is performed independently on each block in shared memory. For the regions which do not fit into shared memory, the computation is done on a CPU. While performance is greatly improved compared to the naive approach on a CPU, the granularity within each block is coarse. The computation of various sized sub-regions creates imbalanced workloads on each thread.

In this work, we focus on the parallel strategies for improving the LRT computation for the 1EXP family. We use a different strategy to ensure workload on each "parallel unit" (PU)² is balanced. We propose a novel range mapping scheme to transform irregular shaped region space to a contiguously regular shaped region space with regards to iterate all of the regions in spatial grid (G). The fine-grained workload on each PU is produced by partitioning the regular space into different equal sized portions. Furthermore, a dynamic pre-computing scheme from our previous work(4) based on the Inclusive/Exclusive principle is presented for 1EXP. Four pre-computed data sets corresponding to four corners of the data grid are generated to reduce the cost of querying the intermediate statistics of each region to $\mathcal{O}(1)$.

Overall, the **contributions** in our work are:

- A novel range mapping scheme is proposed to provide the fine-grained parallelism for LRT computation.
- A dynamic pre-computation scheme is presented for fast computing.
- A k -best reduction strategy is presented for accumulating distributed results on each "PU" and forms the final top- k regions at the end.
- The algorithms are implemented on various parallel architectures and corresponding performances are studied.

The rest of the paper is structured as follows. In Section 2, we provide background materials on LRT computation and its variation on 1EXP family. Motivations are presented in Section 3. In Section 4, we present a novel range mapping scheme for 1EXP LRT computation for multi-dimensional data grid. In Section 5, we explain how we use Inclusive/Exclusive rule and dynamic programming to speed up the enumeration and processing of regions measurements. To produce the final top- k regions, each parallel portion generates top k results and k -best reduction is done on CPU. The proof is provided in Section 6. The details of the implementation on Multi-core, GPGPU and EC2 cloud cluster are presented in Section 7. In Section 8, we evaluate experiments results on these different architectures and discussion is given. Related work is provided in Section 9. We give our conclusion in Section 10.

2 | BACKGROUND

We provide a brief but self-contained introduction for using LRT to find anomalous regions in a spatial setting. The regions are mapped onto a spatial grid G . Given a data set X , an assumed model distribution $f(X, \theta)$, a null hypothesis $H_0 : \theta \in \Theta_0$ and an alternate hypothesis $H_1 : \theta \in \Theta - \Theta_0$, LRT is the ratio

$$\lambda = \frac{\sup_{\Theta_0} \{L(\theta|X)|H_0\}}{\sup_{\Theta} \{L(\theta|X)|H_1\}} \quad (1)$$

where $L()$ is the likelihood function and θ is a set of parameters for the distribution (1). In a spatial setting, the null hypothesis is that the data in a region R (that is currently being tested) and its complement (denoted as \bar{R}) are governed by the same parameters. Thus if a region R is anomalous then the alternate hypothesis will most likely be a better fit and the denominator of λ will have a higher value for the maximum likelihood estimator of θ . A remarkable fact about λ is that under mild regularity conditions, the asymptotic distribution of $\Lambda \equiv -2\log\lambda$ follows a χ_k^2 distribution with k degrees of freedom, where k is the number of free parameters³. Thus regions whose Λ value falls in the tail of the χ^2 distribution are likely to be anomalous (1).

¹The experiment results were reported in 2009.

²We define the (computational) granularity of a parallel architecture as the largest "computational unit" that should run sequentially on an "application computing unit". It refers to "block/thread" for GPGPU, "core" for multi-core and "process" for cloud pc-cluster

³If the χ^2 distribution is not applicable then Monte Carlo simulation can be used to ascertain the p -value.

The one-parameter exponential family (1EXP) simplifies the computation of the LRT statistic. The distribution of a random variable $x \in X$ belongs to a one-parameter exponential family (8) (denoted by $x \sim 1EXP(\theta, \phi, T, B_e, a)$) if it has a probability density given by

$$f(x; \theta) = C(x, \phi) \exp((\theta T(x) - B_e(\theta))/a(\phi)) \quad (2)$$

where $T(\cdot)$ is some measurable function, $a(\phi)$ is a function of some known scale parameter ($\phi > 0$), θ is an unknown parameter (called the natural parameter), $B_e(\cdot)$ is a strictly convex function and $C(\cdot)$ is some constant value. Note that the support of $x : f(x; \theta) > 0$ is independent of θ .

Theorem 1. (8): Let data set $X_R (R \in G)$ be independently distributed with $x_R \sim 1EXP(\theta, \phi, T, B_e, a)$. The log-likelihood ratio test statistic for testing $H_0 : \theta_R = \theta_{\bar{R}}$ versus $H_1 : \theta_R \neq \theta_{\bar{R}}$ is given by:

$$\Lambda = m_R g_e(G \frac{m_R}{b_R}) - \frac{b_R}{G} B_e(g_e(G \frac{m_R}{b_R})) + (1 - m_R) g_e(G \frac{1-m_R}{1-b_R}) - \frac{(1-b_R)}{G} B_e(g_e(G \frac{1-m_R}{1-b_R})) \quad (3)$$

Parameter m_R is the fraction measurement of Region R in total and b_R is the fraction of baseline measure of Region R in total. Correspondingly, $1 - m_R$ is the fraction measurement of Region \bar{R} in total and $1 - b_R$ is the fraction of baseline measure of Region \bar{R} in total. Parameters m_R and b_R are important measurements to calculate the statistic of 1EXP. More information about the functions g_e , G and B_e can be found in (8). For example, if we assume the counts m_R in a region R follow a Poisson distribution with baseline b and intensity λ , then a random variable $x \sim \text{Poisson}(\lambda\mu)$ is a member of 1EXP with $T(x) = x/\mu$, $\Phi = 1/\mu$, $a(\Phi) = \Phi$, $\theta = \log(\lambda)$, $B_e\theta = \exp(\eta)$, $g_e(x) = \log(x)$. For any regions R and \bar{R} , m_R and $m_{\bar{R}}$ are independently Poisson distributed with a mean of $\{\exp(\theta_R)b_R\}$ and $\{\exp(\theta_{\bar{R}})b_{\bar{R}}\}$, respectively. Then, $b_R = \frac{b_R}{b_R + b_{\bar{R}}}$ and $m_R = \frac{m_R}{m_R + m_{\bar{R}}}$. The log-likelihood ratio is calculated by: $c(m_R \log(\frac{m_R}{b_R}) + (1 - m_R) \log(\frac{1-m_R}{1-b_R}))$ (c.f. (8)). The closed-form formula for LRT generalizes to the 1EXP family of distributions (8).

3 | MOTIVATIONS

3.1 | Parallel Enumeration Schemes

In this section, we demonstrate the two different parallel strategies for performing LRT statistics in a one-dimensional data grid by example. The strategies are (i) Overlapping Enumeration, and (ii) Non-overlapping Enumeration. The Overlapping Enumeration strategy has been implemented in (4). In the example, we simulate the architectural environment of (4), and illustrate its drawbacks. The non-overlapping Enumeration strategy, called Range Mapping scheme, will be presented in this paper to overcome the limitations of the Overlapping Enumeration strategy. For simplicity, we will describe the strategies on an abstract parallel machine not detailing its implementation on a concrete parallel platform. The abstract parallel machine has a fixed number of parallel units (PU) whose memory for each PU is finite.

EXAMPLE: Assume we have a one-dimensional spatial data grid with five data points, denoted as $\{O_0, O_1, O_2, O_3, O_4\}$. We would like to compute the LRT of each interval for the one-dimensional spatial data grid. To find the outlier, we enumerate all intervals that can be spawned over the one-dimensional data grid by performing LRT calculation explained in above section 2. We express $l_{(i,j)}$ as an interval between point O_i and O_j , where $(0 \leq i < j \leq n - 1)$. There are $\frac{4 \cdot (4+1)}{2} = 2 \times 5 = 10$ intervals in total, i.e., number of pairs (i, j) for which $(0 \leq i < j \leq 4)$ holds. Figure 1 a visualizes some intervals among above five data points.

To illustrate how to compute LRT for each interval in this example, we assume each interval independently follows poisson distribution and the calculation of LRT is given in last part of section 2 of this paper. The formula is repeated here: $c(m_l \log(\frac{m_l}{b_l}) + (1 - m_l) \log(\frac{1-m_l}{1-b_l}))$, where c is a constant factor and l denotes interval (c.f. (8)). Another detailed similar example in two-dimensional grid is provided in (9). Therefore, to compute LRT value of an interval in the formula, m_l (i.e. success count) and b_l (i.e. population count) are required. Table 2 provides sample value of (m_l, b_l) for each independent interval. To compute the LRT of an interval, there are two different kind of calculations: (a) compute LRT of a single independent interval l , (b) compute LRT of an interval l , which is union of multiple non-overlapped intervals $l = l_1 \cup l_2 \cup \dots \cup l_t$. For example, to obtain the LRT value of a single independent interval $l_{(0,0)}$, the computation is: $l_{rt(0,0)} = 2 \cdot \log(\frac{2}{10}) + (1 - 2) \log(\frac{1-2}{1-10}) = -1.0$. To obtain a larger interval interval $l_{(0,1)}$, which is the union of interval $l_{(0,0)}$ and $l_{(1,1)}$, (m_l, b_l) needs to be aggregated first. Since the data here follows poisson distribution, we know from above that $m_{l(0,1)} = m_{l(0,0)} + m_{l(1,1)} = 2 + 2 = 4$ and $b_{l(0,1)} = b_{l(0,0)} + b_{l(1,1)} = 10 + 10 = 20$. Thus $l_{rt(0,1)} = 4 \cdot \log(\frac{4}{20}) + (1 - 4) \log(\frac{1-4}{1-20}) = -0.9$. Similarly, the LRT of each sub-intervals can be obtained using one of the two approaches.

From above, we know the computation of LRT for each interval. Now we move to present the strategies of parallelizing LRT computation in a data grid. We assume the parameters m_l and b_l of each independent single interval takes up 4 bytes of data storage, respectively. Let's assume that each parallel unit of the abstract parallel machine has at most 16 bytes of data storage. Hence, each parallel unit is limited to compute at most two intervals.

TABLE 1 m_R and b_R statistics for each independent interval $l(i, j)$

Interval	m_R	b_R
(0,0)	2	10
(1,1)	2	10
(2,2)	15	10
(3,3)	3	10

TABLE 2 coefficient a_i and b_i

a_n	a_{n-1}	...	a_0
a	b_n	...	b_0

Overlapping Enumeration Parallel Strategy

Since each parallel unit of the parallel machine can only maximally store and process 16 bytes of data and the statistics values of each single independent interval takes up 8 bytes (i.e. 4 bytes of m_l + 4 bytes of b_l = 8 bytes), therefore the maximum number of interval can be processed is $\frac{PU_{max_allowed_bytes}}{bytes_of_interval} = \frac{16}{8} = 2$ for a single parallel unit at once. In the overlapping strategy (4), each PU_i is associated to k continuous intervals. For this example, each PU enumerates every two continuous overlapping intervals (i.e., the interval (i, j) and the interval $(i, j + 1)$) by a single parallel unit. The LRT s of the remaining intervals will be sequentially processed and merged by the CPU (c.f. (4)). The reasons are: (a) any interval with length $|l| > k$ ($k=2$ in this example) cannot be fitted to a PU due to lack of memory of parallel unit; (b) every two non-continuous single intervals processed on a PU cannot process intervals optimally. On the other hand, if PU store k continuous single intervals, it can process all of the $\frac{k \cdot (k+1)}{2}$ sub-intervals on a single PU. Table 3 illustrates the allocation of continuous intervals to parallel units, and the allocation of the rest intervals to the CPU. For example, PU_0 is assigned the work to compute the LRT for the intervals $\{l_{(0,0)}, l_{(0,1)}\}$, and PU_1 is assigned for computing the LRT for intervals $\{l_{(1,1)}, l_{(1,2)}\}$. Figure 1 a illustrates the enumeration of some intervals for those 5 data points by applying over-lapping strategy.

The aforementioned approach has a few drawbacks: First, non-continuous intervals cannot be processed by parallel units. Hence, there are around $n \cdot \binom{n}{k}$ ($k>2$ in above example) intervals that will be processed sequentially by the CPU. Second, some PU s may have fewer number of intervals as a consequence. For example, PU_4 has only one interval (i.e. $\{l_{(3,3)}\}$). Hence, the overlapping enumeration parallel strategy causes an unbalanced workload among PU s. Third, for each PU , each interval is assigned to a thread. The lengths of intervals are different and this can further creates different workloads among different threads while accumulating statistics for computing LRT . To overcome the drawbacks of this approach, the Non-Overlapping Enumeration parallel strategy is proposed.

Non-Overlapping Enumeration Parallel Strategy

Our new scheme employs a pre-computation strategy are for computing LRT on the abstract parallel machine. Moreover, Horner scheme is used to allocate certain interval to certain PU . The Horner scheme is used to describe elements in a finite discrete multi-dimensional space to a single number and vice versa. Examples are conversion of numbers in different number systems etc.

If we map each interval $l_{i,j}$ as a point (i, j) and plot these points onto two-dimensional space, a triangular shape is formed. In Figure 1 b, all of the 10 intervals forms a triangle shape enclosed by points $(0, 0)$, $(3, 3)$, $(0, 3)$, (i.e. intervals: $l_{(0,0)}, l_{(3,3)}, l_{(0,3)}$). We notice that the total number of intervals is the product of $(n + 1)$ and $\frac{(n)}{2}$. A rectangular shape can be formed by this product: $[width, height] = [n + 1, \frac{n}{2}]$. In this example, $\{(0, 0), (0, 1), (1, 4), (0, 4)\}$ forms a rectangle R_1 and it has the exact same number of points as the formed triangular shape R_2 . Therefore, each point in this rectangle can be mapped to one point in the triangle space. This means each interval in the one-dimensional grid has one-to-one mapping from triangular space to rectangular space (Please see Table 4)

To allocate an interval $l_{(i,j)}$ to a certain PU_y , horner scheme can be used for mapping y to (i, j) and vice versa. To show how horner scheme works, we denote: $f(x) = a_n x^n + \dots + a_1 x + a_0$ (10), where $f(x)$ is described in a polynomial form with coefficients (a_0, a_1, \dots, a_n) within certain field. Such $f(x)$ can be re-written in the form: $f(x) = (x - a)g(x) + r$, where the incomplete fraction is $g(x) = b_{n-1}x^{n-1} + \dots + b_0$ and r is the remainder. See the table 2. For example, in the form of $f(x) = a_0 + x(a_1 + x(a_2 + \dots + x(a_{n-1} + a_n x)))$, the incomplete fraction $a_1 + x(a_2 + \dots + x(a_{n-1} + a_n x))$ and the remainder a_0 can be obtained by given x . To illustrate in more detail, if giving $y \in 0, 1, 3, 4, 5$ in one dimension to map to (i, j) , where $0 \leq i, j < n$, in two dimension, we denote $y = i \cdot n + j$. Using horner scheme, we can derive the fraction i and remainder j uniquely by computing $i = y/n$ and $j = y \bmod n$. Therefore, for a certain PU_y , where $y \in \{0, \frac{n \cdot (n+1)}{2}\}$ a corresponding point (i, j) can be obtained by: $i = y/n$ and $j = y \bmod n$.

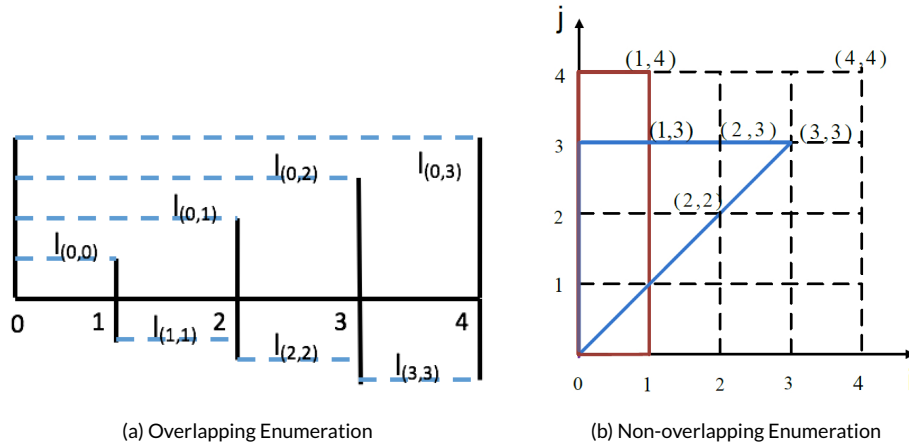


FIGURE 1 Parallel Enumeration Schemes

There are a few advantages: (1) This solves the workload imbalance among *PU*s. Each *PU* processes same amount of intervals. In the rectangular space R_1 , we can assign the same amount of intervals (i.e. points in this example) to each *PU* by dividing the entire space equally. (2) By using a dynamic programming scheme in section 3.2 to pre-compute subsets of *LRT* statistic value and the computation of each thread becomes $\mathcal{O}(1)$, which will solve the workload imbalance issue among threads due to the *LRT* computation on different sized sub-grid for each thread.

TABLE 3 Example: PU assignment

Grid	Intervals on PU		Intervals on CPU
	$interval_1$	$interval_2$	$intervals$
PU_1	$l_{(0,0)}$	$l_{(0,1)}$	$l_{(0,2)}$
PU_2	$l_{(1,1)}$	$l_{(1,2)}$	$l_{(0,3)}$
PU_3	$l_{(2,2)}$	$l_{(2,3)}$	$l_{(1,3)}$
PU_4	$l_{(3,3)}$		

3.2 | Dynamic Pre-computation

From section 2, we know the log-likelihood statistic (*LRT*) computation on 1EXP family is simplified to aggregate the statistic values from a given region R (denoted as \sum). The fraction of the total from actual measurement and baseline measurement m_R, b_R are obtained based on $\sum R$ without direct computation of the complement of R (i.e. \bar{R}). After collecting the aggregated statistic values (m_R, b_R) , theorem 1 is applied to get the *LRT* value of region R .

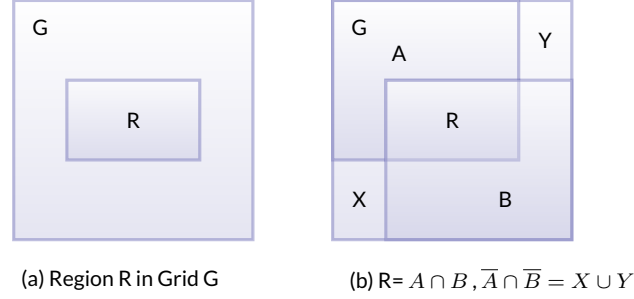
Consider Figure 2 (a) showing a rectangular area R embedded in a grid G . Instead of counting the number of elements in R directly, we express set R as set intersections of two sets A and B as shown in Figure 2 (b). Set A is a rectangular region that starts in the upper left corner of the grid and ends at the lower right corner of R . Set B is a rectangular region that starts at the upper left corner of R and ends at the lower right corner of the grid. Hence, $R = A \cap B$. We denote the region from the lower left corner of G to the lower left corner of R by X and the region from the upper right corner of R to the upper right corner of G by Y .

By applying De Morgan's law and inclusion/exclusion principle, the query of counting the number of elements of region $R(i_1, j_1, i_2, j_2)$, where (i_1, j_1) is the upper left corner and (i_2, j_2) is the lower right corner is expressed by (see Proof in appendix):

$$|R(x_1, y_1, x_2, y_2)| = |A(x_2, y_2)| + |B(x_1, y_1)| + |X(x_1, y_2)| + |Y(x_2, y_1)| - |G| \quad (4)$$

TABLE 4 Interval Transformation of Non-Overlapping Enumeration

$(0,0) \rightarrow (0,0)$	$(0,1) \rightarrow (0,1)$	$(0,2) \rightarrow (0,2)$
$(0,3) \rightarrow (0,3)$	$(0,4) \rightarrow (3,3)$	$(1,4) \rightarrow (2,2)$
$(1,3) \rightarrow (2,3)$	$(1,0) \rightarrow (1,1)$	$(1,1) \rightarrow (1,2)$
$(1,2) \rightarrow (1,3)$		

**FIGURE 2** Set Relations for Region Problem

From Equation 4, we can see that a query time of $\mathcal{O}(1)$ can be achieved by pre-computing statistics of sets A, B, X , and Y for all possible regions in G . Since one of the corner of A, B, X , and Y is fixed, we can pre-compute the cardinalities of these sets in tables of size $\mathcal{O}(n^2)$.

This pre-computation scheme has been implemented in (4) and we will give more details and related algorithms in this paper.

4 | RANGE MAPPING SCHEME

In this section, we study how parallelism helps to enumerate all of the rectangular regions (R) in a spatial grid (G). We will explain this scheme in one-dimensional, two-dimensional, and k -dimensional data-grids.

From section 3.1, we know that if all of the pairwise intervals between n data points, denoted as $l_{(i,j)}$, are plotted as point (i, j) onto two-dimensional coordinate system, they form triangular shape. Each pair of (i, j) in triangular-shaped space can be transformed to pair (i', j') in rectangular-shaped space. After transformation, it enables the whole rectangular space to be partitioned into equal portions. Each portion consists of same amount of pairs and is distributed onto different “parallel computing component” (PU) to balance workload and facilitate parallelization. We name this transforming scheme as range mapping scheme. This scheme can be directly extended to two dimensional spatial grid and multi-dimensional grid.

Figure 3 a plots out all of the intervals $l_{(i,j)}$ in two-dimensional space and we can see that these points (i, j) forms a triangle. Figure 3 b shows the rectangular-shaped space after transformation.

To assign an interval $l_{(i,j)}$ to a PU_y , Horner scheme discussed above is applied (10). For example, we can compute: $i = y/n$ and $j = y \bmod n$, where $y \in (0, \frac{n \cdot (n+1)}{2})$, $y \in (0, n)$. Then transform (i, j) from rectangular space to (i', j') to triangle space. We give details in the following:

4.1 | One-Dimensional Mapping

For one-dimensional range mapping scheme, we assume there are n data points, therefore having $\frac{n \cdot (n+1)}{2}$ intervals. There are two cases in one-dimensional grid: (1) The grid size n is even (2) The grid size n is odd.

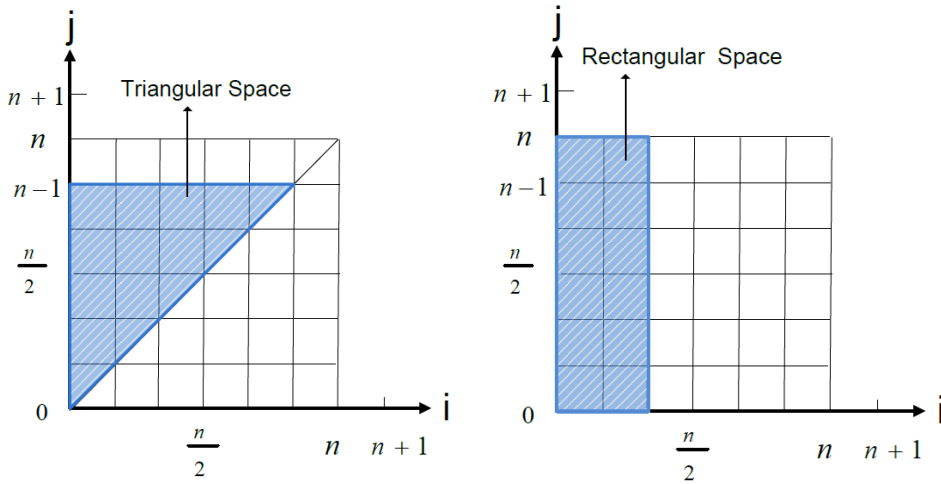


FIGURE 3 1-d Interval Transformation from $[n, n]$ to $[(n+1)/2, n]$

Given the following information:

$$N = \{0, \dots, n-1\} \quad (5)$$

$$R = \{(i, j) \in [0, \dots, \lfloor \frac{n+1}{2} \rfloor - 1] \times [0, \dots, n]\} \quad (6)$$

$$T = \{(i', j') \in [0, \dots, n] \times [0, \dots, n] | i' \leq j'\} \quad (7)$$

$$M = \{0, \dots, \lfloor \frac{n+1}{2} \rfloor \cdot (n+1) - 1\} \quad (8)$$

$$(9)$$

where N is the set of data points which partitions one-dimensional grid to intervals and total number of these points is n . R is the rectangular space formed by $\{[0, \dots, \lfloor \frac{n+1}{2} \rfloor - 1] \times [0, \dots, n]\}$ and each point (i, j) (i.e. interval) in this space is assigned to a certain PU_y . T is the triangular space formed by original intervals, where each interval is $l(i', j')$ ($0 < i' < j' \leq n$). M is total number of intervals for computing LRT .

To have a generalized form of the mapping scheme when grid size n is even or odd, we extend the calculation of total intervals from $\frac{n+1}{2} \cdot n$ to $\lfloor \frac{n+1}{2} \rfloor \cdot (n+1)$. When n is even, we know $\lfloor \frac{n+1}{2} \rfloor = \frac{n}{2}$. When n is odd, we know $\lfloor \frac{n+1}{2} \rfloor = \frac{n+1}{2}$. The generalized computation of number of total intervals in our range mapping scheme is $\frac{n+1}{2} \cdot (n+1)$ no matter grid size n is even or odd, this produces $\frac{n+1}{2}$ more intervals with enlarged set when n is odd but have exact calculation when n is even.

(1) The grid size n is even: A mapping Function $\phi : M \rightarrow R \rightarrow T$ is composed of two functions: $\phi_1 : M \rightarrow R$ and $\phi_2 : R \rightarrow T$. Firstly, ϕ_1 transforms the index value of $y \in M$ of a certain PU_y to a pair $(i, j) \in R$ in rectangular space. Secondly, ϕ_2 transforms the pair $(i, j) \in R$ from rectangular space to a pair $(i', j') \in T$ to triangular space.

Definition 1. IR Mapping Function

$$\phi_1 : M \rightarrow R$$

$$y \mapsto (i, j)$$

where $i = y/(n+1)$ and $j = y \bmod (n+1)$.

IR (Index to rectangular) mapping function assigns a point (i, j) in rectangular space R to y of PU_y by applying inverse horner scheme, where $y \in \{0, \dots, \lfloor \frac{n+1}{2} \rfloor \cdot (n+1) - 1\}$. Point (i, j) is computed by $i = y/(n+1)$ and $j = y \bmod (n+1)$, where y is the index of a particular PU and constant value of n is the total number of data points in one-dimensional grid.

Lemma 1. Mapping function ϕ_1 transforms $y \in M$ (i.e. $y \in \{0, \dots, \lfloor \frac{n+1}{2} \rfloor \cdot n\}$) to $(i, j) \in R$, where $i \in \{0, \dots, \frac{n}{2} - 1\}$ and $j \in \{0, \dots, n\}$.

When n is even, we know $\lfloor \frac{n+1}{2} \rfloor = \frac{n}{2}$. Since $y \in M$ and size of M is equal to $\frac{n}{2} \cdot n$, it is obviously seen that $i \in \{0, \dots, \frac{n}{2} - 1\}$ when $i = y/(n+1)$. Similarly, $j \in \{0, \dots, n\}$ when $j = y \bmod (n+1)$.

Lemma 2. Mapping function $\phi_1 : M \rightarrow R, y \mapsto (i, j)$ is bijective.

In one-dimension grid, for each PU_y , we can denote $y = i \cdot n + j$, where $y \in \{0, \dots, \frac{n \cdot (n+1)}{2}\}$ and $0 \leq i, j < n$. From the horner scheme and the quotient-remainder theorem, we know lemma 1 produces one-to-one mapping from y to (i, j) . At the same time, given (i, j) , y can be uniquely constructed as $y = i \cdot n + j$. Therefore, $y \mapsto (i, j)$ is bijective.

Definition 2. RT Mapping Function

$$\phi_2 : R \rightarrow T \quad (10)$$

$$(i, j) \mapsto (i', j') = \begin{cases} (i, i+j) & \text{if } (i+j) < n \\ (n-i-1, 2n-i-j-1) & \text{otherwise} \end{cases} \quad (11)$$

RT (rectangular to triangular) mapping function transforms point (i, j) in rectangular space to point (i', j') in triangular space shown in above equation. This transformation is achieved by satisfying the constraints: $(i, j) \in \{[0, \dots, \lfloor (n+1)/2 \rfloor] \times [0, \dots, n]\}$ and $0 \leq i' \leq j' < n$. To fulfill this constraint, line $i+j = n$ divide the rectangular space into two sub-spaces: (a) $i+j < n$, (b) $i+j \geq n$. In (a), transformed point (i', j') is calculated as $(i, i+j)$. In (b), transformed point (i', j') is calculated as $(n-i-1, (n-i-1) + (n-j))$ (i.e. $(n-i-1, 2n-i-j-1)$). It is clearly seen that (i', j') satisfies $0 \leq i' \leq j' < n$ after transformation. Figure 4 visualizes the mapping.

Lemma 3. Mapping function ϕ_2 transform $(i, j) \in R$ to $(i', j') \in T$, where $i' \in \{0, \dots, n\}$ and $j' \in \{0, \dots, n\}$.

It is true that $i < n$ and $n-i-1 < n$. Also when $(i+j) < n$ and $j' = i+j$, we can get $i' \leq j' < n$. When $(i+j) \geq n$, we get $i' \leq j' = 2n-i-j-1 = (n-i) + (n-j-1) < n$. Therefore, $i' \in \{0, \dots, n\}$, $j' \in \{0, \dots, n\}$ and $i' \leq j'$.

Lemma 4. Mapping function $\phi_2 : R \rightarrow T, y \mapsto (i, j)$ is bijective.

From above definition, we see that each (i, j) uniquely maps to (i', j') . We get $i' = i, j' = i+j$ when $i+j < n$. And we get $i' = n-i-1, j' = 2n-i-j-1$ when $i+j \geq n$. Correspondingly, given (i', j') , (i, j) can be uniquely obtained by $i = i', j = j' - i'$ or $i = n-i'-1, j = i' + n-j'$. Therefore, $y \mapsto (i, j)$ is bijective.

Corollary 1. The function $\phi : M \rightarrow R \rightarrow T, y \mapsto (i', j')$ is bijective.

Since $M \rightarrow R$ and $R \rightarrow T$ are bijective respectively, this means $y \mapsto (i, j)$ and $(i, j) \mapsto (i', j')$ are bijective. Therefore $M \rightarrow R \rightarrow T, y \mapsto (i', j')$ is bijective.

(2) The grid size n is odd: Function $\phi : M \rightarrow R \rightarrow T$ is composed of two mapping functions: $\phi_1 : M \rightarrow R$ and $\phi_2 : R \rightarrow T$. Firstly, ϕ_1 transforms the index value of $y \in M$ of a certain PU_y to a pair $(i, j) \in R$ in rectangular space. Secondly, ϕ_2 transforms $(i, j) \in R$ from rectangular space to a pair $(i', j') \in T$ in triangular space.

Definition 3. IR Mapping Function

$$\phi_1 : M \rightarrow R \quad (12)$$

$$y \mapsto (i, j) \quad (13)$$

where $i = y/(n+1)$ and $j = y \bmod (n+1)$.

IR (Index to rectangular) mapping function assigns a point (i, j) in rectangular space R to y of PU_y by applying inverse horner scheme, where $y \in \{0, \dots, \lfloor \frac{n+1}{2} \rfloor \cdot (n+1) - 1\}$. Point (i, j) is computed by $i = y/(n+1)$ and $j = y \bmod (n+1)$, where y is the index of a particular PU and constant value of n is the total number of data points in one-dimensional grid. Since n is odd in this case, there has $\frac{n+1}{2}$ more data points (intervals) in the extended form than the original total number of intervals and they won't be processed.

Lemma 5. Mapping function ϕ_1 transforms $y \in M$ to $(i, j) \in R$, where $i \in \{0, \dots, \frac{n+1}{2} - 1\}$ and $j \in \{0, \dots, n\}$.

When n is odd, we know $\lfloor \frac{n+1}{2} \rfloor = \frac{n+1}{2}$. Since $y \in M$ and size of M is extended to $\frac{n+1}{2} \cdot (n+1)$, we can derive that $i \in \{0, \dots, \frac{n+1}{2} - 1\}$ when $i = y/(n+1)$. Similarly, $j \in (0, n)$ when $j = y \bmod (n+1)$.

Lemma 6. Mapping function $\phi_1 : M \rightarrow R, y \mapsto (i, j)$ is bijective.

In one-dimension grid, for each PU_y , we can denote $y = i \cdot (n+1) + j$, where $y \in (0, \frac{(n+1) \cdot (n+1)}{2})$ and $0 \leq i, j < n$. From the horner scheme and the quotient-remainder theorem, we know there is unique mapping from y to (i, j) . Correspondingly, given (i, j) , y can be uniquely constructed as $y = i \cdot (n+1) + j$. Therefore, $y \mapsto (i, j)$ is bijective.

Definition 4. Mapping Function

$$\phi_2 : R \rightarrow T \quad (14)$$

$$(i, j) \mapsto (i', j') = \begin{cases} (i, i+j) & \text{if } (i+j) < n \\ (n-i-1, 2n-i-j-1) & \text{if } (i+1) \leq \frac{(n+1)}{2} \end{cases} \quad (15)$$

RT (rectangular to triangular) mapping function transforms point (i, j) in rectangular space to point (i', j') in triangular space shown in above equation. This transformation is achieved by satisfying the constraints: $(i, j) \in \{[0, \dots, \lfloor (n+1)/2 \rfloor] \times [0, \dots, n+1]\}$ and $0 \leq i' \leq j' < n$. To fulfill this constraint, line $i+j = n$ divide the rectangular space into two sub-spaces: (a) $i+j < n$, (b) $i+j \geq n$. Furthermore, when n is odd, we have $\frac{(n+1)}{2}$ more points than needed and thus the rectangular shape is enlarged, the constraint of $(i+1) \leq \frac{(n+1)}{2}$ is applied to limit the unnecessary transformation of points in rectangular space to triangular space. To explain this, when line $i+j = n$ intersection with line $(i+1) = \frac{(n+1)}{2}$, we can get $j = n-i = n - \frac{(n+1)}{2} + 1 = \frac{(n+1)}{2}$. Hence when $i+j < n$, the unnecessary points above the intersection point on the line of $(i+1) = \frac{(n+1)}{2}$ are filtered out. In (a), transformed point (i', j') is calculated as $(i, i+j)$. In (b), transformed point (i', j') is calculated as $(n-i-1, (n-i-1) + (n-j))$ (i.e. $(n-i-1, 2n-i-j-1)$). This mapping strategy shows that (i', j') satisfies $0 \leq i' \leq j' < n$ after transformation. Figure 4 b visualizes the mapping.

Lemma 7. Mapping function $\phi_2 : R \rightarrow T$ transform $(i, j) \in R$ to $(i', j') \in T$, where $i' \in \{0, \dots, n\}$ and $j' \in \{0, \dots, n\}$ in triangular space.

It is true that $i < n$ and $n-i-1 < n$. Also when $(i+j) < n$ and $j' = i+j$, we can get $i' \leq j' < n$. When $(i+j) \geq n$, we get $i' \leq j' = 2n-i-j-1 = (n-i) + (n-j-1) < n$. Moreover, the constraint of $(i+1) \leq \frac{(n+1)}{2}$ is applied to limit the unnecessary transformation of points when n is odd. Therefore, $i' \in \{0, \dots, n\}$, $j' \in \{0, \dots, n\}$ and $i' \leq j'$ in triangular space.

Lemma 8. Mapping function $\phi_2 : R \rightarrow T, (i, j) \mapsto (i', j')$ is injective.

From above mapping strategy, we see that each (i', j') has single mapping to (i, j) and some unnecessary (i, j) are not processed due to R is enlarged. There has $i' = i, j' = i+j$ when $i+j < n$. And there has $i' = n-i-1, j' = 2n-i-j-1$ when $i+j \geq n$. There is no overlapping mapped point (i', j') based on the constraint of $i+j < n$ and $(i+1) \leq \frac{(n+1)}{2}$. Correspondingly, given (i', j') , (i, j) can be obtained by $i = i', j = j' - i'$ or $i = n - i' - 1, j = i' + n - j'$. However, due to R is enlarged space, $(i, j) \mapsto (i', j')$ is injective when n is odd.

Corollary 2. Mapping function $\phi : M \rightarrow R \rightarrow T, y \mapsto (i', j')$ is injective.

From above, we know $M \rightarrow R$ (i.e. $y \mapsto (i, j)$) is bijective and $R \rightarrow T$ (i.e. $(i, j) \mapsto (i', j')$) is injective. Therefore $M \rightarrow R \rightarrow T, y \mapsto (i', j')$ is injective.

Figure 4 a and Figure 4 b show the solutions and implementation is shown in Algorithm 1.

Example: To illustrate the transformation, Figure 5 a and 5 b show small examples on how to transform the intervals when having data points $\{0, 1, 2, 3, 4\}$ and $\{0, 1, 2, 3, 4, 5\}$ respectively. For data point $(i, j) \in \{0, 1, 2, 3, 4\}$, the total intervals is 10. By plotting out all of them in Figure 5 a, the triangular shaped space is bounded by $\{(0, 0), (3, 3), (0, 3)\}$ and there are exactly 10 points in total. The rectangular shaped space is bounded by $\{(0, 0), (1, 0), (1, 4), (0, 4)\}$ and total number of points is exactly 10. Similarly, for data point $(i, j) \in \{0, 1, 2, 3, 4, 5\}$, the triangular shaped space is bounded by $\{(0, 0), (4, 4), (0, 4)\}$ and total number of points is 15. The rectangular shaped space is bounded by $\{(0, 0), (2, 0), (2, 5), (0, 5)\}$ and total number of points is 15. Table 5 shows the transformations.

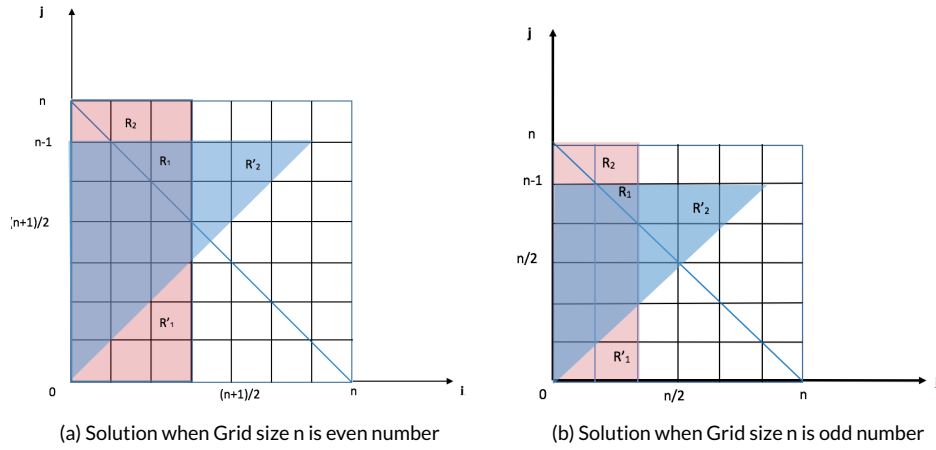


FIGURE 4 1-d Interval Transformation from $[n, n]$ to $[\lfloor (n+1)/2, n+1]$

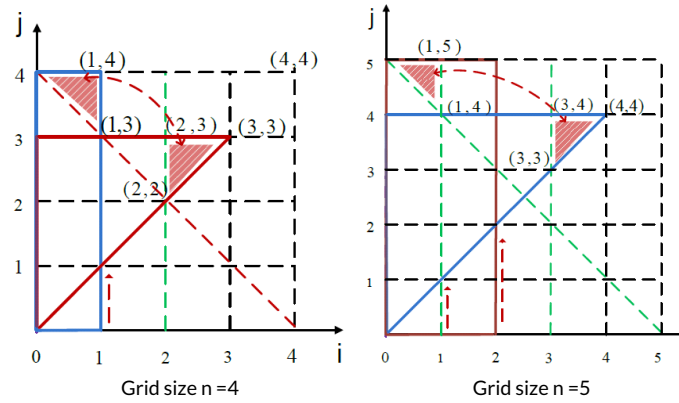


FIGURE 5 Example: 1-d Interval Transformation

$(0,0) \rightarrow (0,0)$	$(0,1) \rightarrow (0,1)$	$(0,2) \rightarrow (0,2)$
$(0,3) \rightarrow (0,3)$	$(0,4) \rightarrow (3,3)$	$(1,4) \rightarrow (2,2)$
$(1,3) \rightarrow (2,3)$	$(1,0) \rightarrow (1,1)$	$(1,1) \rightarrow (1,2)$
$(1,2) \rightarrow (1,3)$		

Grid (4,4)

$(0,0) \rightarrow (0,0)$	$(0,1) \rightarrow (0,1)$	$(0,2) \rightarrow (0,2)$
$(0,3) \rightarrow (0,3)$	$(0,4) \rightarrow (0,4)$	$(0,5) \rightarrow (4,4)$
$(1,0) \rightarrow (1,1)$	$(1,1) \rightarrow (1,2)$	$(1,2) \rightarrow (1,3)$
$(1,3) \rightarrow (1,4)$	$(1,4) \rightarrow (3,4)$	$(1,5) \rightarrow (3,3)$
$(2,0) \rightarrow (2,2)$	$(2,1) \rightarrow (2,3)$	$(2,3) \rightarrow (2,4)$

Grid (5,5)

TABLE 5 Example: Interval Transformation for even/odd number of points

4.2 | Two-Dimensional Mapping

Two-dimensional mapping is extended from one-dimensional mapping directly.

Given:

$$N_{1,2} = \{0..n_{1,2} - 1\} \quad (16)$$

$$R_{1,2} = \{0..\lfloor \frac{n_{1,2}+1}{2} \rfloor - 1\} \times \{0..n_{1,2}\} \quad (17)$$

$$T_{1,2} = \{(a_{1,2}, b_{1,2}) \in N_{1,2}^2 | a_i \leq b_{1,2}\} \quad (18)$$

$$M_{1,2} = \{0..\lfloor \frac{n_{1,2}+1}{2} \rfloor \times (n_{1,2} + 1) - 1\} \quad (19)$$

$$(20)$$

where N_1, N_2 are coordinate range, R_1, R_2 are rectangular space, T_1, T_2 are triangular space and M_1, M_2 are number range in two-dimension.

Mapping function $\phi^2 : M^2 \rightarrow R^2 \rightarrow T^2$ does transformation. Horner scheme is applied to get two-dimensional tuple (x_1, x_2) . Then ϕ^2 is used to transform (x_1, x_2) to (i_1, j_1, i_2, j_2) .

4.3 | K-Dimensional Mapping

Similarly, in K-dimension, by given:

$$N_k = \{0..n_k - 1\} \quad (21)$$

$$R_k = \{0..\lfloor \frac{n_k+1}{2} \rfloor - 1\} \times \{0..n_k\} \quad (22)$$

$$T_k = \{(a_k, b_k) \in N_k^2 | a_i \leq b_k\} \quad (23)$$

$$M_k = \{0..\lfloor \frac{n_k+1}{2} \rfloor \times (n_k + 1) - 1\} \quad (24)$$

$$(25)$$

where k is the number of dimension transformed, N_k is coordinate range, R_k is rectangular space, T_k is triangular space and M_k is number range in K-dimension.

Mapping function $\phi^k : M^k \rightarrow R^k \rightarrow T^k$ performs transformation. Horner's scheme is used to transform a number in M^k to a k-dimensional tuple (x_1, \dots, x_k) . Then mappings ϕ^k performs the translation from each element x_i to $(i_1, \dots, k, j_1, \dots, k)$.

5 | INCLUSIVE/EXCLUSIVE PRE-COMPUTATION SCHEME

Motivated by section 3.2, we now present a novel and unique approach, based on the Inclusive/Exclusive principle and dynamic programming to accumulate the statistic of a region R . We build a table in time $\mathcal{O}(n^2)$ and then compute the statistic of any region R in $\mathcal{O}(1)$ time.

To obtain a query time of $\mathcal{O}(1)$, we need to pre-compute sets A, B, X , and Y for all possible regions in G . Since one of the corner is fixed we can pre-compute the cardinalities of these sets in tables of size $\mathcal{O}(n^2)$.

To obtain the tables for A, B, X , and Y , we employ dynamic programming. The dependency and statistic counts' propagation of rows and columns for these tables are shown in Figure 6 a, 6 b, 6 c and 6 d. For example, the table for A can be computed using the following recurrence relationship:

$$|A(i, j)| = |A(i, j-1)| + |A(i-1, j)| - |A(i-1, j-1)| + |G(i, j)| \quad (26)$$

where $|G(i, j)|$ ⁴ counts whether there is an element in the cell location (i, j) . The first element and the first column and row need to be populated (initialized) so that all cardinalities of A can be computed. The counts in the remaining rows and columns are accumulated through the dependency of the previous row and column. Figure 6 a shows the computation of set A and the implementations of it is listed in Algorithm 2 (see the proofs and the rest implementation of set B, X, Y in Appendix.)

Similarly, the computation of set B, X, Y is listed as the following:

⁴(i,j) represents (x_i, x_j) for simplicity.

Algorithm 1 Parallel Range Mapping in 1-d array

Input: n data points

Output: mapping all the intervals from rectangular to triangular space

```

1: //The interval (i,j) in rectangular space is transformed to interval (i',j') in triangular space
2: for i ← 0 to (n+1)/2 do
3:   for j ← 0 to n do
4:     if j < (n - i) then
5:       i' ← i
6:       j' ← (i+j)
7:     else
8:       if 2(i + 1) < (n + 1) then
9:         i' ← (n-i+1)
10:        j' ← (n-j+i')
11:      end if
12:    end if
13:    LRT computation for interval (i,j)
14:  end for
15: end for

```

$$|B(i, j)| = |B(i + 1, j)| + |B(i, j + 1)| - |B(i + 1, j + 1)| + |G(i, j)| \quad (27)$$

$$|X(i, j)| = |X(i, j + 1)| + |X(i - 1, j)| - |X(i - 1, j + 1)| + |G(i, j)| \quad (28)$$

$$|Y(i, j)| = |Y(i + 1, j)| + |Y(i, j - 1)| - |Y(i + 1, j - 1)| + |G(i, j)| \quad (29)$$

Due to the high dependency among rows and columns, pre-computing is hard to parallelise and the computation is very fast on a CPU. In our work, the pre-computation of set A , B , X , Y is done on a CPU.

6 | K-BEST REDUCTION SCHEME

To find top- k^5 anomalous rectangular regions, a heap with maximum size of k is built from each “parallel computing component” (PU). A further reduction strategy is applied on these k heaps to get final k -best regions.

A LRT value set, denoted as $s = \{s_1, s_2, \dots, s_i, \dots, s_n\}$, is generated from n rectangular regions set R . $\{R\}$ is divided into t equal portions: $\{p_1, p_2, \dots, p_i, \dots, p_t\}$, where $1 \leq t \leq n$. Each portion is processed in parallel and a k -best result with heap structure is generated correspondingly. The k -best result from each portion is $p_{1k}, p_{2k}, \dots, p_{ik}, \dots, p_{tk}$.

We denote the process of finding k -best using heap sort as $h(\cdot)$. We also denote the k -best result from original data grid is $\{s_{r0}, s_{r1}, \dots, s_{rk}\}$. And the k -best value from each parallel portion p_i is $\{s_{i0}, s_{i1}, \dots, s_{ik}\}$. And we assume the k -best value are in ascending order. For example, $s_{r0} \leq s_{r1}, \dots, \leq s_{rk}, s_{i0} \leq s_{i1}, \dots, \leq s_{ik}$.

Lemma 9. *KBestReduction* : The k -best LRT values from the value set s is equal to the k -best values obtained by performing k -best reduction from each parallel portion: $h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n) = h(h(p_1) \cup h(p_2) \cup \dots \cup h(p_n))$.

We give proof by contradiction:

⁵Top- k and k -best are exchangeable terms.

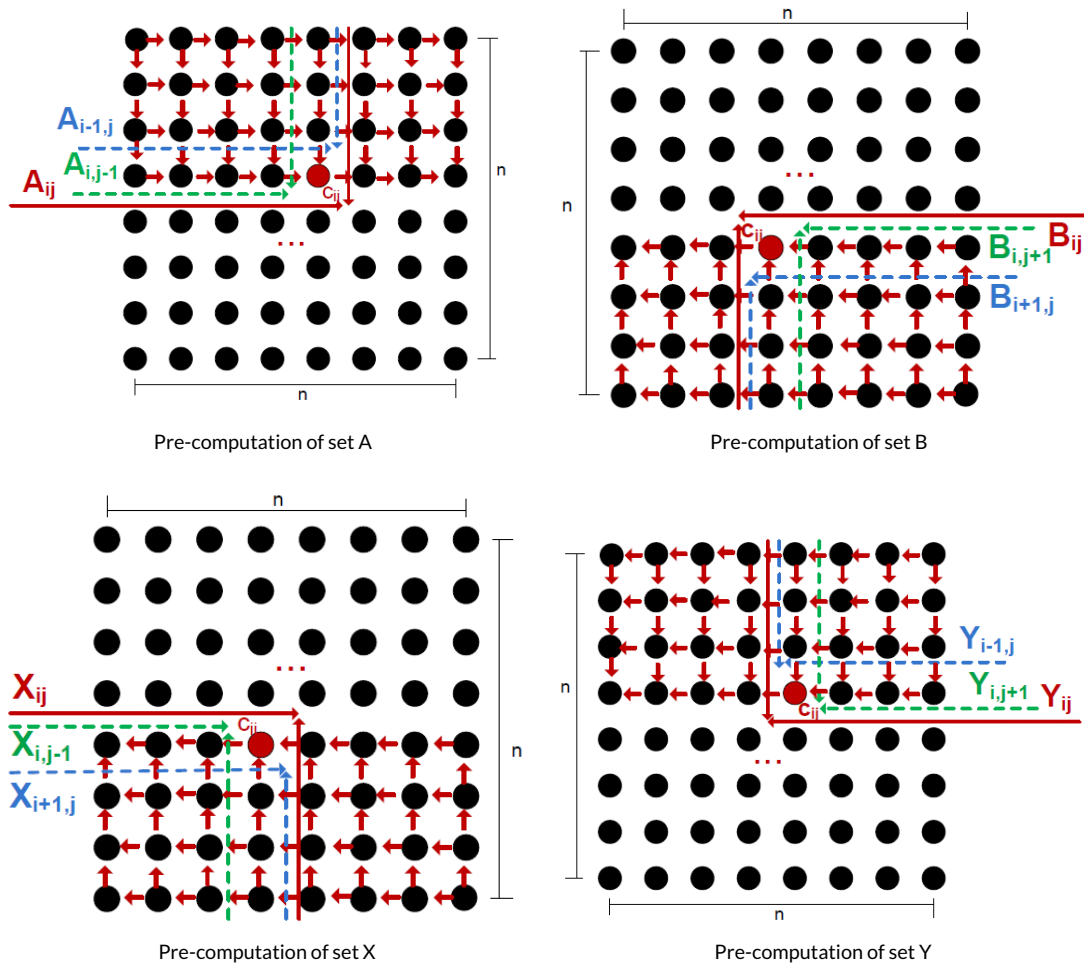


FIGURE 6 pre-computation of set A, B, X and Y

$$(1) \forall x \in h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n)$$

We have

$$x \in (h(p_1) \cup h(p_2) \cup \dots \cup h(p_n)), \quad (30)$$

$$x \in h(h(p_1) \cup h(p_2) \cup \dots \cup h(p_n)) \quad (31)$$

Because the total number of k -best values: $|(h(p_1) \cup h(p_2) \cup \dots \cup h(p_n))| \geq k$,

and $x \notin h(p_1) \cup h(p_2) \cup \dots \cup h(p_n) \rightarrow x \notin h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n)$.

and x is one of the topk value from all the results

Therefore, $x \in h((h(p_1) \cup h(p_2) \cup \dots \cup h(p_n)))$

$$(2) \forall x \in h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n)$$

We have

$$x \in h(p_1) \cup h(p_2) \cup \dots \cup h(p_n), \quad (32)$$

$$x \in h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n) \quad (33)$$

Because $h(h(p_1) \cup h(p_2) \cup \dots \cup h(p_n)) \subseteq h(p_1) \cup h(p_2) \cup \dots \cup h(p_n) \subseteq h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n)$

Therefore, $x \in h(p_1 \cup p_2 \cup \dots \cup p_i \cup \dots \cup p_n)$

From (1) and (2), So the k -best from the whole value set s is same as the reduction from the k -best results from each processes. We implemented *minheap* and *maxheap* to store the k -best results for each parallel threads/process.

Algorithm 2 Inclusive/Exclusive Pre-computation for Set A

Input: data grid (G)

Output: accumulated counts $A(i, j)$

```

1: //Initialize first element  $A(0, 0)$ 
2:  $A(0, 0) \leftarrow G(0, 0)$ 
3: //accumulation of remaining elements in first row
4: for  $j \leftarrow 1$  to  $n$  do
5:    $A(0, j) \leftarrow G(0, j) + A(0, j - 1)$ 
6: end for
7: //accumulation of remaining elements in first column
8: for  $i \leftarrow 1$  to  $n$  do
9:    $A(i, 0) \leftarrow G(i, 0) + A(i - 1, 0)$ 
10: end for
11: //accumulation of all the elements in remaining rows and columns
12: for  $k \leftarrow 1$  to  $n$  do
13:   for  $i \leftarrow k$  to  $n$  do
14:      $A(i, k) \leftarrow G(i, n + k) + A(i - 1, k) + A(i, k - 1) - A(i - 1, k - 1)$ 
15:   end for
16:   for  $j \leftarrow k$  to  $n$  do
17:      $A(k, j) \leftarrow G(k, n + j) + A(k, j - 1) + A(k - 1, j) - A(k - 1, j - 1)$ 
18:   end for
19: end for

```

7 | IMPLEMENTATIONS ON PARALLEL AND DISTRIBUTED ARCHITECTURE

To detect anomalous regions in spatial/spatial-temporal grid, for brute-force approach, all the contiguous and rectangular areas are searched and LRT computation is performed over each of them. The regions within top-k LRT statistic values are treated as potential outliers. However, it is very time-consuming to identify the outliers.

To improve performance, pre-computation method is presented based on inclusive/exclusive scheme. Furthermore, we devise range mapping scheme to transform the total rectangular regions from triangular shaped space to rectangular shaped space. This mapping have a contiguous space without gap to enable the workload perfectly partitioned into same amount for each PU.

We implemented the naive approach, inclusive/exclusive approach on a single CPU and range mapping scheme on different parallel and distributed architectures: Multi-core, GPGPU and EC2 Cloud Cluster in two-dimensional grid (n, n) .⁶

7.1 | Naive Approach

The implementation of naive approach of LRT computation on a two-dimensional grid (n, n) is shown in algorithm 3. We can see the computation complexity of brute-force searching is $\mathcal{O}(n^4)$ and the worst case of the computation of single LRT for a region R is $\mathcal{O}(n^2)$.

7.2 | Inclusive/Exclusive Approach

Inclusive/Exclusive scheme in section 3 is applied on the whole spatial grid (G) to generate four pre-computed data set. For any given region R , retrieving the intermediate statistic value is $\mathcal{O}(1)$ time. The overall enumeration cost is reduced in a sequential version. The k -best heap structure is built to keep global top-k regions. The implementation is shown in algorithm 4.

⁶Please see source code (11).

Algorithm 3 Naive top-k LRT searchInput: data grid $(G(n,n))$, k Output: top k anomalous regions

```

1: //Search each rectangle  $R(x1,y1,x2,y2)$  in  $G(n,n)$ 
2: for  $x1 \leftarrow 0$  to  $n$  do
3:   for  $y1 \leftarrow 0$  to  $n$  do
4:     for  $x2 \leftarrow x1$  to  $n$  do
5:       for  $y2 \leftarrow y1$  to  $n$  do
6:          $score \leftarrow lrt(x1,y1,x2,y2)$ 
7:       end for
8:     end for
9:   end for
10: end for
11: Sorting the scores and return top-k regions  $R$ 

```

Algorithm 4 Inclusive/Exclusive top-k LRT searchInput: data grid $(G(n,n))$, k Output: top k anomalous regions

```

1: //Inclusive/Exclusive pre-computation
2:  $prefix\_sums \leftarrow compute\_prefix\_sums(A,B,X,Y)$ 
3: //Search each rectangle  $R(x1,y1,x2,y2)$  in  $G(n,n)$ 
4: for  $x1 \leftarrow 0$  to  $n$  do
5:   for  $y1 \leftarrow 0$  to  $n$  do
6:     for  $x2 \leftarrow x1$  to  $n$  do
7:       for  $y2 \leftarrow y1$  to  $n$  do
8:          $intermediate\ statistics \leftarrow prefix\_sums(x1,y1,x2,y2)$ 
9:          $score \leftarrow lrt(x1,y1,x2,y2,intermediate\ statistics)$ 
10:        //keep kbest regions
11:         $kbest \leftarrow heap(R,G)$ 
12:      end for
13:    end for
14:  end for
15: end for

```

7.3 | Multi-core Approach

Multi-core computers, consisting two or more processors working together on a single integrated circuit, have produced breakthrough performance. They allow faster execution of applications by taking advantage of parallelism, or the ability to work on multiple problems simultaneously. The default mechanism for running parallel programs on multi-core processors is to run a separate process on each core. Threads enable a lighter weight approach than processes. A standardized programming interface called POSIX (Portable Operating System Interface for Unix) was developed to support multiple architecture and provide the capabilities of threads. Implementations adhering to this standard are refereed as Pthreads. Pthreads provide task and data parallelism and flexibility to the programmers.

There are two key advantages associated with LRT computation work on a multi-core architecture with Pthreads. Firstly, for a 1EXP family data grid, LRT computation of each rectangular region is independent. This provides the task parallelism. Secondly, the non-gap ranging mapping scheme clearly enables us to divide the workload to each thread equally, improving scaling.

Algorithm 5 LRT implementation on *Multicore* Architecture: Main Part

Input: data grid (G), likelihood function (f)

Output: top k anomalous regions

```

1: //declare threads array
2: pthread_t ← thread[num_threads]
3: thread_param ← param[num_threads]
4: // do pre-computation
5: prefix_sums ← compute_prefix_sums()
6: // total workload be partitioned
7: size ← ((n + 1)(n + 1))2 / 4

8: // compute the workload of each thread
10: stride ← size/num_threads

12: // initialize thread parameters and spawn threads
13: for i ← 0 to num_threads do
14:   thread_param[i].start ← start
15:   thread_param[i].end ← start + stride
16:   thread_param[i].prefix ← prefix_sums
17:   create_pthread(thread_param)
18: end for
19: parallel processing (see the algorithm of parallel part)
20: //get kbest regions
21: kbest ← heap(kbest[0,...,num_threads])

```

Algorithm 6 LRT implementation on Multi-core: Parallel Part

Input: prefix-sum, thread id (i) portion index start, end

Output: kbest regions processed by thread i

```

1: // thread i process regions from (start,end)
2: for portion index ← start to end do
3:   LRT computation for regions obtained from index
4:   // keep kbest for current thread i
5:   kbest ← heap(i)
6: end for

```

The implementation of multi-core version is shown in algorithm 5 and algorithm 6. In the main program, each thread is assigned with the same workload and same size of range. It is equal to: total number of regions to be searched divides by the defined number of threads. The range mapping scheme enables each thread to process in a non-gap rectangular range. Otherwise, it is hard to provide each thread same size of range. Each thread performs LRT computation on its own part and keeps k -best regions using min, max heap structure. After all threads finishes computation, the heap function is applied to all the k -best results and get the final k -best regions.

7.4 | GPU/Multi-GPU Approach

The General Purpose Graphic Processing Unit (GPGPU) architecture allows graphic card to be used as general purpose parallel computing device. We briefly describe how the NVIDIA's Compute Unified Device Architecture (CUDA) framework supports massively parallel computing (12).

Algorithm 7 GPUkernel

Input: a spatial grid $G(n,n)$, number of blocks (bx,by) , number of threads per block (tx,ty)

Output: top k anomalous regions processed by each thread

```

1: //total regions to be enumerated
2:  $total\_workload = (n(n+1)/2)^2$ 
3: //compute the workload for each thread
4:  $tx\_size \leftarrow (n(n+1)/2 + bx \cdot tx - 1) / (bx \cdot tx)$ 
5:  $ty\_size \leftarrow (n(n+1)/2 + by \cdot ty - 1) / (by \cdot ty)$ 
6: //LRT computation of each thread
7: for  $i \leftarrow 0$  to  $tx\_size$  do
8:   for  $j \leftarrow 0$  to  $ty\_size$  do
9:     //get region coordinates from range mapping scheme
10:     $(x_1, y_1, x_2, y_2) \leftarrow reverse\_range\_mapping(i, j)$ 
11:    //perform LRT computation on region R
12:     $score \leftarrow lrt(R(x_1, y_1, x_2, y_2))$ 
13:    //keep kbest for current thread
14:     $kbest \leftarrow heap(current\ thread)$ 
15:   end for
16: end for

```

A CUDAProgram consists of one or more phases that are executed on either the host (CPU) or the device (GPU). The host code executed on CPU exhibits little or no data parallelism and the device code (called kernel) executed on GPU side exhibits high amount of data parallelism. The kernel uses large number of threads to exploit data parallelism. The CUDATHreads are extremely light weight and require very few clock cycles to be generated and scheduled on the GPU. The key programming challenge is to map the computation and expressing them onto an abstract model consisting of grids, blocks and threads. The kernel is then executed by a grid of thread blocks. Blocks execute concurrently among multiprocessors while threads in a given block execute concurrently within a single multiprocessor. Only threads within a block can cooperate with each other.

To exploit CUDA for the LRT computation, all of regions are mapped from triangular range shape onto rectangular range shape. The rectangular range is divided into different non-overlap blocks with same size. Each block has a number of threads to process a number of rectangular regions. For fast computation, each block searches all of the rectangular regions which can be fitted into shared memory. Each thread is assigned to compute a subset of rectangle. To avoid transferring all the results back CPU, parallel reduction is performed. Each thread keeps a k -best heap. Furthermore, keeping the total k -best heaps not arbitrary, it might lead to another kernel for parallel reduction. By defining block number and thread number for different size of data grid, each thread is responsible for varied number of rectangles in different kernel. We did the implementation on single GPGPU and multi-GPGPU and measured the performance. The implementation of single GPGPU is shows in algorithm 9 . For multi-GPGPU implementation, we divide the whole workload equally to each GPGPU. Final k -best regions are obtained after each device return back all the k -best heaps.

7.5 | Amazon EC2 Cloud Cluster

Cloud computing offers a highly scalable infrastructure for high performance computing. Amazon Elastic Compute Cloud (Amazon EC2) enables ?compute? in the cloud. It is possible to get a set of computing instances on demand without requiring a lot of maintenance and financial resources a common cluster would need. We present our parallel implementation for cloud computing using MPI. The workload is equally assigned to each process based on our range mapping scheme and each process performs LRT computation on each rectangle and produces k -best regions with heap structure using reduce function. The implementation is shown in algorithm 8.

Algorithm 8 MPI main program**Input:** a spatial grid $G(n,n)$, number of processes np , top-k $kbest$ **Output:** top-k anomalous rectangular regions

```

1: //initialize
2: MPI_init()
3: //broadcast grid information to all nodes
4: MPI_Bcast(n,n,g)
5: total_workload = (n(n + 1)/2)2
6: //compute workload for each process
7: stride ← (total_workload + np - 1)/np
8: start ← processid · stride
9: end ← start + stride
10: //parallel LRT computation
11: Rectangle r ← compute(g, start, end)
12: //store kbest results for each process
13: kbest ← heap(r)
14: // reduce operation
15: MPI_reduce(kbest)
16: //root process
17: if processid == 0 then
18:   kbest ← qsort(reduced_kbest)
19: end if
20: MPI_Finalize()

```

8 | EXPERIMENTS AND ANALYSIS

8.1 | Performance Analysis

We have designed and implemented a set of experiments to show and validate the performance speedup on the above different architectures. The experiments are performed on a Poisson distribution model and a randomly generated anomalous region was introduced for verification in synthetic data sets. Answers to the following questions were sought:

- What are the performance gains of naive versus Inclusive/Exclusive approach?
- How is the multi-core scaling of the LRT computation on Multi-core architecture?
- What are the performance gains of GPGPU approaches versus Inclusive/Exclusive approach?
- How is the MPI scaling of LRT computation on PC-cluster?
- How is the computation speed of LRT computation for (n, n) grid on these architectures?

8.1.1 | Naive vs. Inclusive/Exclusive Implementation

We run the naive and inclusive/exclusive implementation on an 8-core E5520 Intel server to compare their performance. Data grid varies from (100, 100) to (1000, 1000). The results are shown in Figure 7 a and Figure 7 b.

From the results in Figure 7 a, we can see that Inclusive/Exclusive approach has a significant speed-up. The number of rectangles per second processed by Inclusive/Exclusive approach is also plotted for various grid sizes in Figure 7 b, it can be seen the result is almost constant, validating that our Inclusive/Exclusive pre-computation approach search complexity is $\mathcal{O}(1)$.

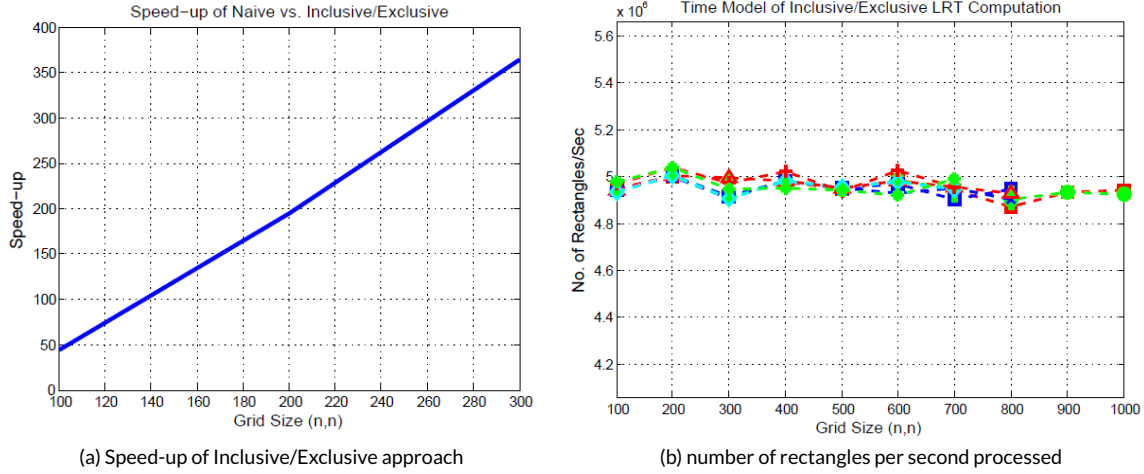


FIGURE 7 Naive vs. Inclusive/Exclusive Approach

8.1.2 | Multi-core

The multi-core experiment is conducted on a 32-core AMD Opteron(tm) Processor 6128 server with 128GB of RAM and a 64-core AMD Opteron(tm) Processor 6378 server with 384GB of RAM. We use data grid with size of (1000,1000) to show the performance scaling. Figure 8 a and 8 c show that the 1EXP-LRT computation scales very well on multi-core architecture. The speedup increases near-linearly with the increase of number of cores (noc) and is consistent with the $\mathcal{O}(n^4/noc)$ running time on each core. A speedup of n on n cores is nearly achieved, indicating near perfect scaling.

The number of rectangles processed per second is plotted against the number of cores in Figure 8 b and 8 d. The results show the linearity of increasing number of rectangles processed per second with increasing number of cores.

8.1.3 | Multi-GPGPU

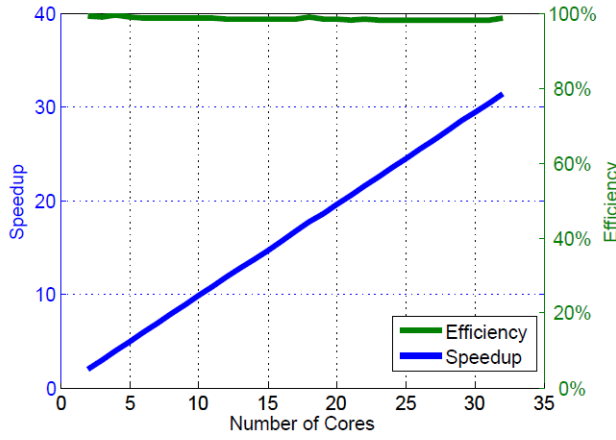
The experiments are conducted on two different GPGPU configurations:

(1) An 8-core E5520 Intel server that is equipped with two GPGPU *Tesla C1060* cards supporting CUDA 4.0. Each GPGPU card has 4GB global memory, 16KB shared memory, 240 cores and 30 multiprocessors. Experiments are conducted to compare the speed-up performance of GPGPUs with exclusive/inclusive cpu approach (see figure 9 a) on various data grid size (n,n). Furthermore, the performance of number of rectangles processed per second on one single GPGPU vs. two GPGPUs are compared (see figure 9 b). The results show that LRT computation on two GPGPUs is around 2 times faster than that on one single GPGPU regardless of the data grid size. And the computation cost of per rectangle is almost constant with the increase of data grid for single GPGPU and two GPGPUs with exclusive/inclusive cpu approach.

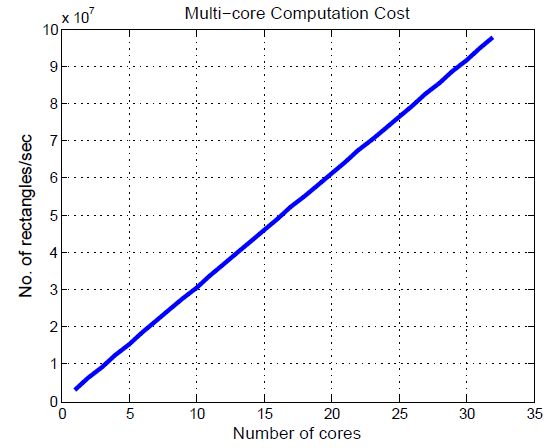
(2) An 8-core server that is equipped with two GPGPU *Tesla K20m* cards supporting CUDA 6.5. Each GPGPU card has 5GB global memory, 64KB shared memory, 2496 cores and 13 multiprocessors. In this architecture, we investigate the speed-up, efficiency and the number of rectangles processed per second with different number of threads on different grid size (see figure 9 c and figure 9 d). Figure 9 c shows the linearly increasing speed-up with increasing number of threads on a given grid size. The efficiency is almost 100%. Figure 9 d shows the linear increasing speed of processing number of rectangles with the increasing of threads on a given grid size.

Given the $((n+1) \times (n+1))/2^2$ regions to be searched for a (n,n) grid and the number of threads in each block is (tx, ty) . The number of blocks (bx, by) changes the LRT computation performance. In our implementation, each region object takes 8byte. To fully utilize the shared memory, each block has $(tx, ty) = (16, 8)$ threads. We vary the value of (bx, by) for the grids with different size to find the optimal block configuration. To maximize performance:

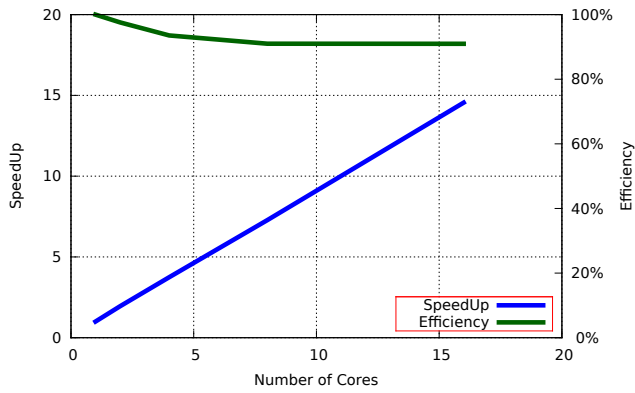
- $Max(blocks) \leq (bx \times by) \leq ((n+1) \times (n+1))/2^2$, otherwise some blocks won't work.
- $((n+1) \times (n+1))/2^2 / (bx \times by) \geq (tx \times ty)$ makes each thread processes at least one region.
- $((n+1) \times (n+1))/2^2 / tx \leq (((n+1) \times (n+1))/2) / ty$ is better for reducing bank conflicts since it retrieves more data by rows.



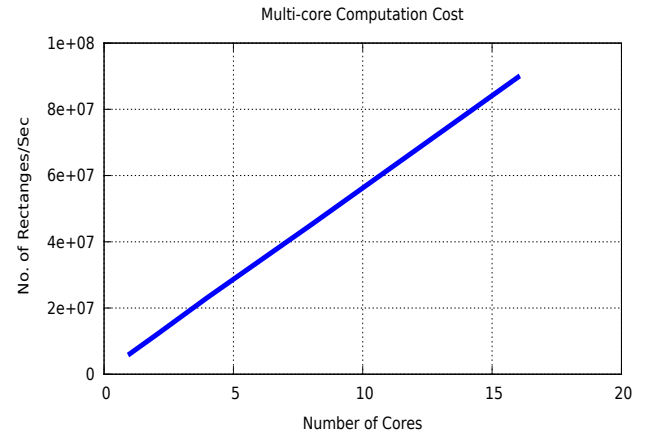
(a) Speed-up and Efficiency of Grid (1000,1000) on Multi-core



(b) The number of rectangles processed per second vs. number of cores on Grid (1000,1000)



(c) Speed-up and Efficiency of Grid (1000,1000) on Multi-core 2



(d) The number of rectangles processed per second vs. number of cores on Grid (1000,1000) on Multi-core2

FIGURE 8 Multi-core Evaluation

TABLE 6 Optimized Block Configuration

<i>Grid</i>	<i>block_x</i>	<i>block_y</i>
(500,500)	128	64
(600,600)	192	86
(700,700)	176	86

If there are too few blocks, each thread processes a quite number of regions and thus performance is degraded. Table 6 gives the optimal block configuration. A speedup of up to two can be achieved by choosing the right block configuration using the slowest run for a given grid size as baseline.

8.1.4 | EC2 Cloud Cluster

We study a cluster composed of 20 *EC2* high-CPU compute nodes and its effect on MPI application scaling. Instances of this family have proportionally more CPU resources than memory (RAM) and are well suited for compute-intensive applications. Figure 10 a shows the nearly linear speedup with the increase of number of processes for different grid size on cloud cluster. The computation speed of rectangles is plotted out in Figure 10 b,

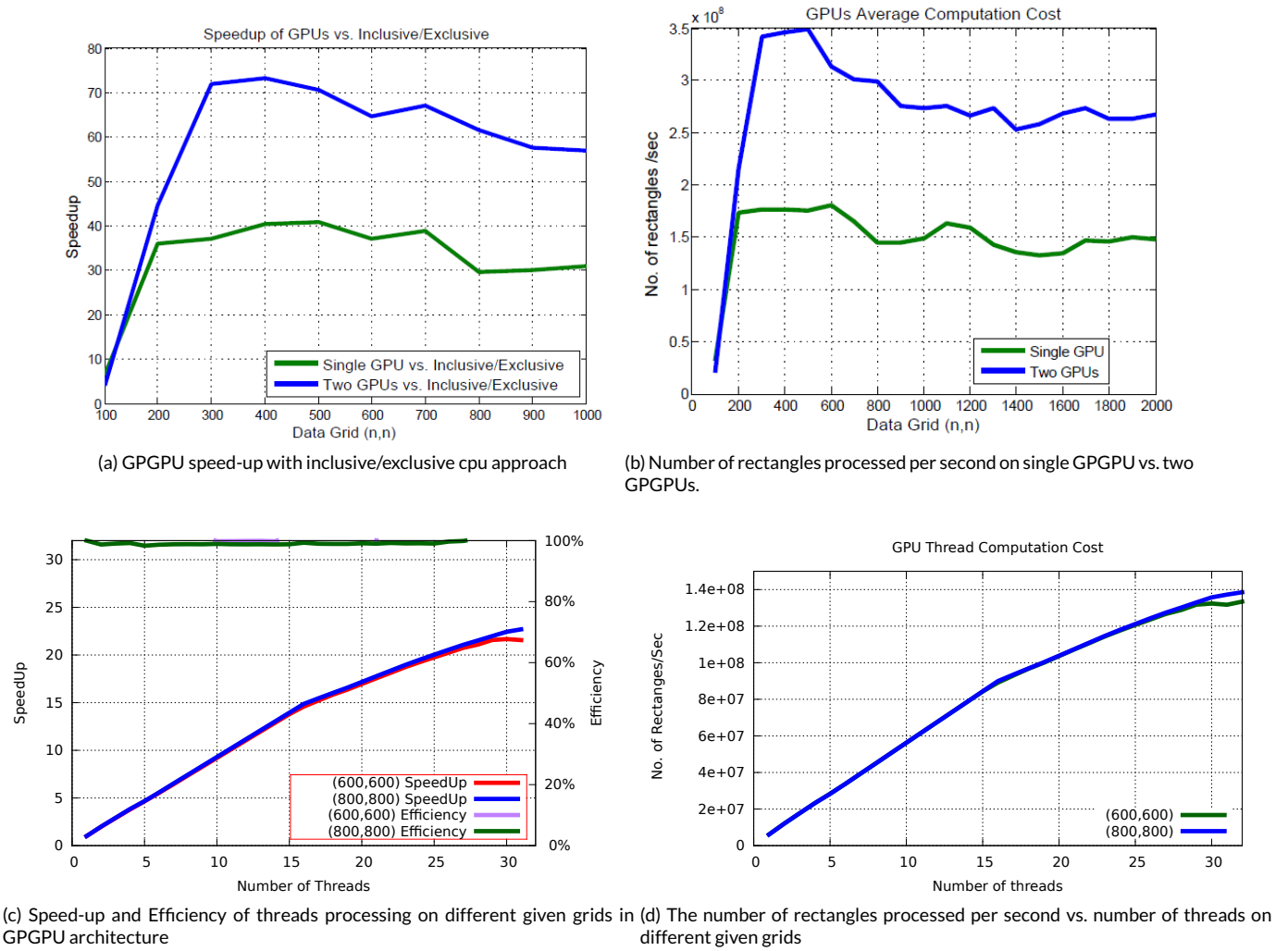


FIGURE 9 GPGPUs Cost

it shows the computation speed is faster with the increase number of processes on a given data grid and also verifies the constant computation speed for different data grid by a given number of processes.

8.2 | Discussion

From the above results, we further analyze of LRT computation on different architectures. We plot the number of rectangles processed per second for data grid (1000, 1000) on the following different architectures: multi-core, GPGPU and EC2 cloud cluster. The results are shown in Figure ?? . The dashed line in the figure is the processing speed of Inclusive/Exclusive approach on single CPU. From the figure, we can see that the GPGPU approach performs much better than multi-core and EC2, which is almost one order of magnitude faster. Furthermore, we can see that, as the number of cores and the number of processes increase, the LRT processing speed is improved on these architectures.

9 | RELATED WORK

Previous attempts to parallelize LRT computation have only achieved limited success. For example, the Spatial Scan Statistic (SSS), which is a special case of LRT for Poisson data, is available as a program under the name SatScan (13). It has been parallelized for multi-core CPU environments

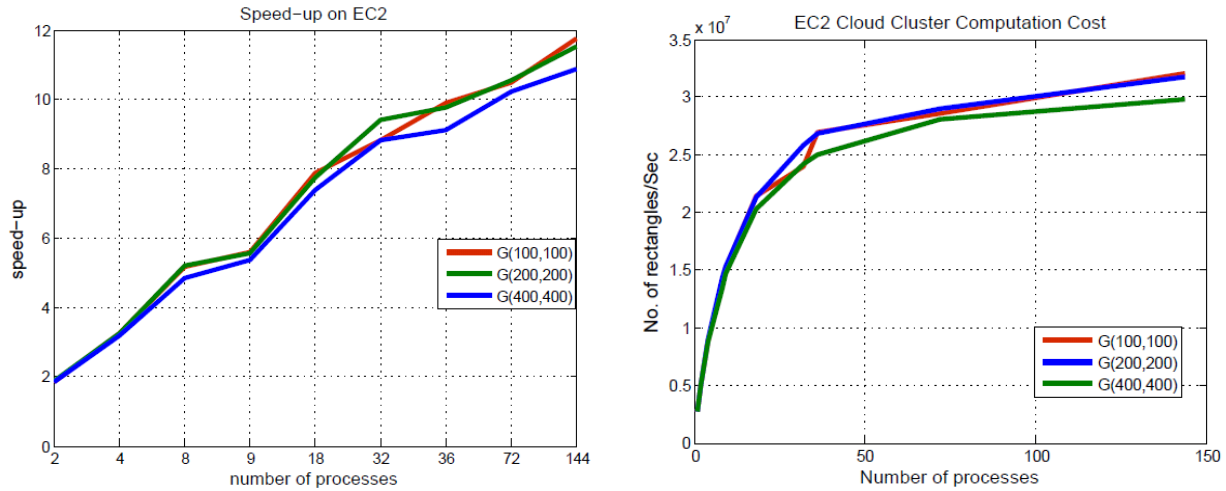


FIGURE 10 Speed-up of LRT computation on EC2 cloud cluster

and its extension for a GPGPU hardware (14) has achieved improved speed up of two over the multi-core baseline. The GPGPU implementation in (14) has proposed loading parts of the data into shared memory but has achieved only a modest speed up. The other attempt of (15) applied their own implementation of a spatial scan statistic program on the GPU to the epidemic disease dataset. This solution is only applicable to its special disease scenario. In each of these cases, we believe there is further room for optimising the algorithms for the parallel architectures by devising the fine-grained parallelism strategies.

Furthermore, all the existing parallel solutions perform LRT tests in a circular or cylindrical way, not in a grid-based scenario. Our parallel solution is different and provides a fully paralleled template for 1EXP-LRT computation in a grid.

10 | CONCLUSION

The Likelihood Ratio Test Statistic (LRT) is a state-of-the-art method for identifying hotspots or anomalous regions in large spatial settings. To speed up the LRT computation for 1EXP family, this paper proposed three ideas: (i) a novel range mapping scheme is proposed to fully enumerate all the regions in a contiguous space. (ii) a dynamic pre-computation algorithm is implemented to reduce the cost of aggregating intermediate statistics. (iii) to save space and improve processing speed, kbest reduction scheme is presented to accumulate distributed results. We did the implementations on different parallel architectures: Multi-core, Multi-GPGPU and EC2 cloud cluster and extensive experiments are done correspondingly. From the results, we see that pre-computation approach has a linear speed-up with the data grid size comparing to the brute-force sequential approach. Then we compare the speed-up on different parallel architectures using pre-computation approach as baseline. The speed-up of these parallel approaches increases near-linearly with the increase of the number of “parallel computing component” on different architectures. In concert, the parallel approaches yield a speed up of nearly four thousand times compared to their sequential counterpart. Further analysis on the processing speed of number of rectangles is given. This provides some recommended information for choosing the right architectures on various factors. Moving the computation of the LRT statistics to the parallel architectures enables the use of this sophisticated method of outlier detection for larger spatial grids than previously reported.

In future, we will apply the range mapping scheme on non-1EXP family distribution using pruning strategy (1). A unified parallel approach will be provided for generalized LRT computation in spatial grids.

SUPPORTING INFORMATION

11 | INCLUSIVE/EXCLUSIVE STATISTICS AGGREGATION

$$|R(x_1, y_1, x_2, y_2)| = |A(x_2, y_2)| + |B(x_1, y_1)| + |X(x_1, y_2)| + |Y(x_2, y_1)| - |G| \quad (34)$$

Proof.

$$|R| = \sum_{i=1}^{x_2} \sum_{j=1}^{y_2} c_{i,j} + \sum_{i=x_1}^n \sum_{j=y_1}^n c_{i,j} + \sum_{i=x_2}^n \sum_{j=1}^{y_1} c_{i,j} + \sum_{i=1}^{x_1} \sum_{j=y_2}^n c_{i,j} \quad (35)$$

$$- \sum_{i=1}^n \sum_{j=1}^n c_{i,j} \quad (36)$$

$$= \sum_{i=1}^{x_1} \sum_{j=1}^{y_2} c_{i,j} + \sum_{i=x_1}^{x_2} (\sum_{j=1}^{y_1} c_{i,j} + \sum_{j=y_1}^{y_2} c_{i,j}) + \sum_{i=x_1}^n \sum_{j=y_1}^n c_{i,j} \quad (37)$$

$$+ \sum_{i=x_2}^n \sum_{j=1}^{y_1} c_{i,j} + \sum_{i=1}^{x_1} \sum_{j=y_2}^n c_{i,j} \quad (38)$$

$$= \sum_{i=1}^{x_1} \sum_{j=1}^n c_{i,j} + \sum_{i=x_1}^{x_2} \sum_{j=1}^{y_1} c_{i,j} + |R| + |R| + \sum_{i=x_1}^{x_2} \sum_{j=y_2}^n c_{i,j} \quad (39)$$

$$+ \sum_{i=x_2}^n \sum_{j=1}^n c_{i,j} - \sum_{i=1}^n \sum_{j=1}^n c_{i,j} \quad (40)$$

$$= \sum_{i=1}^n \sum_{j=1}^n c_{i,j} + |R| - \sum_{i=1}^n \sum_{j=1}^n c_{i,j} \quad (41)$$

$$= |R| \quad (42)$$

□

Definition 5.

$$A(i, j) = \sum_{k=1}^i \sum_{l=1}^j c_{k,l} \quad \text{for all } 1 \leq i, j \leq n. \quad (43)$$

Lemma 10.

$$A(i, j) = A_{i-1,j} + A_{i,j-1} - A_{i-1,j-1} + c_{i,j} \quad \text{for all } 1 < i, j \leq n. \quad (44)$$

Proof.

$$A(i, j) = \sum_{k=1}^{i-1} \sum_{l=1}^j c_{k,l} + a_{i,j-1} - a_{i-1,j-1} + c_{i,j} \quad (45)$$

$$= \sum_{k=1}^{i-1} \left(\sum_{l=1}^{j-1} c_{k,l} + c_{k,j} \right) + a_{i,j-1} - \sum_{k=1}^{i-1} \sum_{l=1}^{j-1} c_{k,l} + c_{i,j} \quad (46)$$

$$= \sum_{k=1}^{i-1} c_{k,j} + c_{i,j} + a_{i,j-1} \quad (47)$$

$$= \sum_{k=1}^i c_{k,j} + a_{i,j-1} \quad (48)$$

$$= \sum_{k=1}^i c_{k,j} + \sum_{k=1}^i \sum_{l=1}^{j-1} c_{k,l} \quad (49)$$

$$= A(i, j) \quad (50)$$

□

Definition 6.

$$X_{i,j} = \sum_{k=i}^n \sum_{l=1}^j c_{k,l} \quad \text{for all } 1 \leq i, j \leq n. \quad (51)$$

Lemma 11.

$$X_{i,j} = X_{i,j-1} + X_{i+1,j} - X_{i+1,j-1} + c_{i,j} \quad \text{for all } 1 < i, j \leq n. \quad (52)$$

Proof.

$$\begin{aligned}
 X_{i,j} &= \sum_{k=i+1}^n \sum_{l=1}^j c_{k,l} + X_{i,j-1} - X_{i+1,j-1} + c_{i,j} \\
 &= \sum_{k=i+1}^n \left(\sum_{l=1}^{j-1} c_{k,l} + c_{k,j} \right) + X_{i,j-1} - \sum_{k=1+1}^n \sum_{l=1}^{j-1} c_{k,l} + c_{i,j} \\
 &= \sum_{k=i+1}^n c_{k,j} + c_{i,j} + X_{i,j-1} \\
 &= \sum_{k=i}^n c_{k,j} + X_{i,j-1} \\
 &= \sum_{k=i}^n c_{k,j} + \sum_{k=i}^n \sum_{l=1}^{j-1} c_{k,l} \\
 &= X_{i,j}
 \end{aligned}$$

□

Definition 7.

$$X_{i,j} = \sum_{k=i}^n \sum_{l=1}^j c_{i,j} \quad \text{for all } 1 \leq i, j \leq n. \quad (53)$$

Lemma 12.

$$X_{i,j} = X_{i,j-1} + X_{i+1,j} - X_{i+1,j-1} + c_{i,j} \quad \text{for all } 1 < i, j \leq n. \quad (54)$$

Proof.

$$\begin{aligned}
 X_{i,j} &= \sum_{k=i+1}^n \sum_{l=1}^j c_{k,l} + X_{i,j-1} - X_{i+1,j-1} + c_{i,j} \\
 &= \sum_{k=i+1}^n \left(\sum_{l=1}^{j-1} c_{k,l} + c_{k,j} \right) + X_{i,j-1} - \sum_{k=1+1}^n \sum_{l=1}^{j-1} c_{k,l} + c_{i,j} \\
 &= \sum_{k=i+1}^n c_{k,j} + c_{i,j} + X_{i,j-1} \\
 &= \sum_{k=i}^n c_{k,j} + X_{i,j-1} \\
 &= \sum_{k=i}^n c_{k,j} + \sum_{k=i}^n \sum_{l=1}^{j-1} c_{k,l} \\
 &= X_{i,j}
 \end{aligned}$$

□

Definition 8.

$$y_{i,j} = \sum_{k=1}^i \sum_{l=j}^n c_{i,j} \quad \text{for all } 1 \leq i, j \leq n. \quad (55)$$

Lemma 13.

$$Y_{i,j} = Y_{i-1,j} + Y_{i,j+1} - Y_{i-1,j+1} + c_{i,j} \quad \text{for all } 1 < i, j \leq n. \quad (56)$$

Algorithm 9 Inclusive/Exclusive Pre-computation for Set B

Input: data grid (G)

Output: accumulated counts $B(i, j)$

```

1: //Initialize first element  $B(n-1, n-1)$ 
2:  $B(n-1, n-1) \leftarrow G(n-1, n-1)$ 
3: //accumulation of remaining elements in last column
4: for  $j \leftarrow (n-1)$  to 1 do
5:    $B(j-1, n-1) \leftarrow G(i-1, n-1) + B(j, n-1)$ 
6: end for
7: //accumulation of remaining elements in last row
8: for  $i \leftarrow (n-1)$  to 1 do
9:    $B(n-1, i-1) \leftarrow G(n-1, i-1) + B(n-1, i)$ 
10: end for
11: //accumulation of all the elements in remaining rows and columns
12: for  $k \leftarrow 1$  to  $n$  do
13:   for  $i \leftarrow (n-1)$  to  $k$  do
14:      $B(i-k, n-1-k) \leftarrow G(i-k, n-1-k) + B(i-k+1, n-1-k) + B(i-k, n-k) - B(i-k+1, n-k)$ 
15:   end for
16:   for  $j \leftarrow (n-1)$  to  $k$  do
17:      $B(n-1-k, j-k) \leftarrow G(n-1-k, j-k) + B(n-1-k, j-k+1) + B(n-k, j-k) - B(n-k, j-k+1)$ 
18:   end for
19: end for

```

Proof.

$$\begin{aligned}
Y_{i,j} &= \sum_{k=1}^{i-1} \sum_{l=j}^n c_{k,l} + Y_{i,j+1} - Y_{i-1,j+1} + c_{i,j} \\
&= \sum_{k=1}^{i-1} \left(\sum_{l=j+1}^n c_{k,l} + c_{k,j} \right) + Y_{i,j+1} - \sum_{k=1}^{i-1} \sum_{l=j+1}^n c_{k,l} + c_{i,j} \\
&= \sum_{k=1}^{i-1} c_{k,j} + c_{i,j} + Y_{i,j+1} \\
&= \sum_{k=1}^i c_{k,j} + Y_{i,j+1} \\
&= \sum_{k=1}^i c_{k,j} + \sum_{k=1}^i \sum_{l=j+1}^n c_{k,l} \\
&= Y_{i,j}
\end{aligned}$$

□

12 | PRE-PROCESSING OF INCLUSIVE/EXCLUSIVE COMPUTATION**References**

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Algorithm 10 Pre-processing of Inclusive/Exclusive Computation for Set X

```

1: //Initialize first column
2: for  $i \leftarrow 1$  to  $n$  do
3:    $X(i,0) \leftarrow 0$ 
4: end for
5: //Initialize last row
6: for  $j \leftarrow 1$  to  $n$  do
7:    $X(n-1,j) \leftarrow 0$ 
8: end for
9: //Iterate over all diagonal elements
10: for  $k \leftarrow 1$  to  $n$  do
11:   //associate columns of diagonal elements
12:   for  $i \leftarrow n-1$  to  $k$  do
13:      $X(i-k,k) \leftarrow G(i-k+1,k-1) + X(i-k+1,k) + X(i-k,k-1) - X(i-k+1,k-1)$ 
14:   end for
15:   // associate columns of diagonal element k
16:   for  $j \leftarrow k$  to  $n$  do
17:      $X(n-1-k,j) \leftarrow G(n-k,j-1) + X(n-1-k,j-1) + X(n-k,j) - X(n-k,j-1)$ 
18:   end for
19: end for

```

Algorithm 11 Inclusive/Exclusive Pre-computation for Set Y

Input: data grid (G)

Output: accumulated counts $Y(i, j)$

```

1: //Initialize last column
2: for  $i \leftarrow 1$  to  $n$  do
3:    $Y(i,n-1) \leftarrow 0$ 
4: end for
5: //Initialize first row
6: for  $j \leftarrow 1$  to  $n$  do
7:    $Y(0,j) \leftarrow 0$ 
8: end for
9: //Iterate over all diagonal elements
10: for  $k \leftarrow 1$  to  $n$  do
11:   //associate columns of diagonal elements
12:   for  $i \leftarrow k$  to  $(n-1)$  do
13:      $Y(i,n-1-k) \leftarrow G(i-1,n-k) + Y(i-1,n-1-k) + Y(i,n-k) - Y(i-1,n-k)$ 
14:   end for
15:   // associate columns of diagonal element k
16:   for  $j \leftarrow (n-1)$  to  $k$  do
17:      $Y(k,n-k) \leftarrow G(k-1,n-k+1) + Y(k,n-k+1) + Y(k-1,n-k) - Y(k-1,n-k+1)$ 
18:   end for
19: end for

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