

# A Compositional Framework for Developing Parallel Programs on Two-Dimensional Arrays

Kento EMOTO ([emoto@ipl.t.u-tokyo.ac.jp](mailto:emoto@ipl.t.u-tokyo.ac.jp))  
, Zhenjiang HU ([hu@mist.i.u-tokyo.ac.jp](mailto:hu@mist.i.u-tokyo.ac.jp))  
, Kazuhiko KAKEHI ([kaz@ipl.t.u-tokyo.ac.jp](mailto:kaz@ipl.t.u-tokyo.ac.jp))  
and Masato TAKEICHI ([takeichi@mist.i.u-tokyo.ac.jp](mailto:takeichi@mist.i.u-tokyo.ac.jp))  
*Department of Mathematical Informatics, Graduate School of Information Science  
and Technology, The University of Tokyo*

**Abstract.** Computations on two-dimensional arrays such as matrices and images are one of the most fundamental and ubiquitous things in computational science and its vast application areas, but development of efficient parallel programs on two-dimensional arrays is known to be hard. In this paper, we propose a compositional framework that supports users, even with little knowledge about parallel machines, to develop both correct and efficient parallel programs on dense two-dimensional arrays systematically. The key feature of our framework is a novel use of the abide-tree representation of two-dimensional arrays. The presentation not only inherits the advantages of tree representations of matrices where recursive blocked algorithms can be defined to achieve better performance, but also supports transformational development of parallel programs and architecture-independent implementation owing to its solid theoretical foundation - the theory of constructive algorithmics.

**Keywords:** Constructive Algorithmics; Skeletal Parallel Programming; Matrix.

## 1. Introduction

Computations on two-dimensional arrays, such as matrix computations, image processing, and relational database managements, are both fundamental and ubiquitous in computational science and its vast application areas <sup>(15,23,34)</sup>. Developing efficient parallel algorithms for these computations is one of the most important topics in many textbooks on parallel programming <sup>(22,36)</sup>. Then, algorithms have been designed and implemented as standard libraries. For example, for matrix computations <sup>(20,40)</sup>, we have the useful libraries like ScaLAPACK<sup>(14)</sup>,

PLAPACK<sup>(1)</sup> and RECSY<sup>(30)</sup>. Though these are useful, there are some limitations when using these libraries to develop parallel programs for manipulating two-dimensional arrays.

- First, the libraries are of low abstraction, and thus difficult to be modified or adapted to solve slightly different problems. For example, we cannot easily change the operators used in matrix multiplications. In fact, the increasing popularity of parallel machines like PC clusters enables more and more users to utilize such parallel computer environments to perform parallel computations of various kinds, which can naturally be slightly different from those the libraries provide. The libraries are of no direct help for the users here, and they have to rewrite or develop the libraries for themselves. However, since (re-)building parallel libraries is much more involved than sequential algorithms due to necessity of considering the synchronization and communication between processors, not everyone can do it easily.
- Second, the libraries are not well structured, and thus hard to be efficiently composed together in the sense that we may need to convert intermediate data structures between two different libraries. Often each library is carefully designed with suitable data structures and algorithms so that it can be efficiently executed on specific parallel architectures. This may, however, prevents us from making efficient use of two libraries developed for two different parallel architectures.

This situation demands a new programming model allowing users to describe parallel computation over two-dimensional arrays in an

easy, efficient, but compositional way. As one promising solution to the demand, *skeletal parallel programming* using the parallel skeleton is known <sup>(11,13,35)</sup>. In this model, users can build parallel programs by composing ready-made components (called *skeletons*) implemented efficiently in parallel for various parallel architectures. This compositional approach has two major advantages: (1) since low-level parallelism is concealed in skeletons, users can obtain a comparatively efficient parallel program without needing detailed techniques of parallel computers and being conscious of parallelism explicitly, (2) since the skeletons are designed for structured programming, they can be efficiently composed to solve various problems.

Much research has been devoted to parallel skeletons on lists, a one-dimensional data structure, and it has been shown <sup>(25,27)</sup> that parallel programming with list skeletons is very powerful since we can describe many problems with these list skeletons. Moreover much research has been done on methods of deriving and optimizing parallel programs with parallel skeletons on lists <sup>(10,21,24)</sup>, and especially about optimization, and there is a library that can automatically optimize a program described by skeletons <sup>(32)</sup>. Similarly, for parallel skeletons on the tree data structure there is research on binary trees <sup>(19,39)</sup>, general trees and derivation of programs on these tree skeletons.

Despite the success of parallel programming with list or tree skeletons, it remains as a big challenge <sup>(33)</sup> to design a skeletal framework for developing parallel programs for manipulating two-dimensional arrays. Generally, a skeleton (compositional) framework for manipulating two-dimensional arrays should consist of the following three parts:

- *a fixed set of parallel skeletons* for manipulating two-dimensional arrays, which cannot only capture fundamental computations on two-dimensional arrays but also be efficiently implemented in parallel for various parallel architectures;
- *a systematic programming methodology*, which can support developing both efficient and correct parallel programs composed by these skeletons; and
- *an automatic optimization mechanism*, which can eliminate inefficiency due to compositional or nested uses of parallel skeletons in parallel programs.

Our idea is to use the theory of constructive algorithmics (also known as *Bird-Meertens Formalism*) <sup>(7,9,38)</sup>, a successful theory for compositional sequential program development. In this theory, aggregate data types are formalized *constructively* as an algebra, and computations on the aggregate data are structured as *recursive* mappings between algebras while enjoying nice algebraic properties for composition with each other.

The key is to formalize two-dimensional arrays constructively so that we can describe computations on them as recursions with maximum (potential) parallelism, allowing implementation to have the maximum freedom to reorder operations for efficiency on parallel architectures. The traditional representations of two-dimensional arrays by nested one-dimensional arrays (row-major or column-major representations) <sup>(29,38)</sup> impose much restriction on the access order of elements. Wise et al. represent a two-dimensional array by a quadtree <sup>(42)</sup> and show that recursive algorithms on quadtree provide better

performance than existing algorithms for some matrix computations (QR factorization <sup>(18)</sup>, LU factorization <sup>(43)</sup>). More examples can be found in the survey paper <sup>(15)</sup>. However, the unique representation of two-dimensional arrays by quadtrees does not capture the whole information a two-dimensional data may have. In contrast, Bird <sup>(7)</sup> represents two-dimensional arrays by dynamic trees allowing restructuring trees when necessary.

In this paper, we propose a compositional framework that allows users, even with little knowledge about parallel machines, to describe safe and efficient parallel computation over two-dimensional arrays easily, and enables derivation and optimization of programs. The main contributions of this paper are summarized as follows.

- We propose a *novel use of the abide-tree representation of two-dimensional arrays* <sup>(7)</sup> in developing *parallel* programs for manipulating two-dimensional arrays, whose importance has not been fully recognized in parallel programming community. Our abide-tree representation of two-dimensional arrays not only inherits the advantages of tree representations of matrices where recursive blocked algorithms can be defined to achieve better performance <sup>(15,18,43)</sup>, but also supports systematic development of parallel programs and architecture independent implementation owing to its solid theoretical foundation - the theory of constructive algorithmics <sup>(7,9,38)</sup>.
- We provide a *strong programming support* for developing both efficient and correct parallel programs on two-dimensional arrays in a highly abstract way (without the need to be concerned with low

level implementation). In our framework (Section 4), programmers can easily build up a complicated parallel system by defining basic components *recursively*, combining components *compositionally*, and improving efficiency *systematically*. The power of our approach can be seen from the nontrivial programming examples of matrix multiplication and QR decomposition <sup>(18)</sup>, and a successful derivation of an involved efficient parallel program for the maximum rectangle sum problem <sup>(2,3,24,41)</sup>.

- We demonstrate an *efficient implementation* of basic computation skeletons (in C++ and MPI) on distributed PC clusters, guaranteeing that programs composed by these parallel skeletons can be efficiently executed in parallel. So far most research focuses on showing that the recursive tree representation of matrices is suitable for parallel computation on shared memory systems <sup>(15,18)</sup>, this work shows that the recursive tree representation is also suitable for distributed memory systems. In fact, our parallel skeletons, being of high abstraction with all potential parallelism, are architecture-independent.

Our framework can be considered as an extension of the quadtree framework of Wise et al. in the sense that it imposes no restriction on the size and the element order of two-dimensional arrays and provides an support of derivation and optimization of programs on two-dimensional arrays.

The rest of this paper is organized as follows. In Section 2, we construct a theory of the abide-trees. Then we design a fixed set of parallel skeletons and give some examples of parallel algorithms on the abide-

trees in Section 3. Section 4 demonstrates systematic development of parallel programs on two-dimensional arrays. Section 5 gives efficient implementations and shows the experimental results. We discuss the related work in Section 6, and make conclusion in Section 7.

An extended version of this paper is available in the technical report (16).

## 2. Theory of Two-Dimensional Arrays

Before explaining our compositional programming framework, we will construct a theory of two-dimensional arrays, the basis of our framework, based on the theory of constructive algorithmics <sup>(7,9,38)</sup>.

Notation in this paper follows that of Haskell <sup>(8)</sup>, a pure functional language that can describe both algorithms and algorithmic transformation concisely. Function application is denoted by a space and the argument may be written without brackets. Thus,  $f a$  means  $f(a)$  in ordinary notation. Functions are curried, i.e. functions take one argument and return a function or a value, and the function application associates to the left. Thus,  $f a b$  means  $(f a) b$ . The function application binds more strongly than any other operator, so  $f a \otimes b$  means  $(f a) \otimes b$ , but not  $f (a \otimes b)$ . Function composition is denoted by  $\circ$ , so  $(f \circ g) x = f (g x)$  from its definition. Binary operators can be used as functions by sectioning as follows:  $a \oplus b = (a \oplus) b = (\oplus b) a = (\oplus) a b$ . For arbitrary binary operator  $\otimes$ , an operator in which the arguments are swapped is denoted by  $\tilde{\otimes}$ . Thus,  $a \tilde{\otimes} b = b \otimes a$ . Two binary operators  $\ll$  and  $\gg$  are defined by  $a \ll b = a$ ,  $a \gg b = b$ . Pairs are Cartesian products of

plural data, written like  $(x, y)$ . A function that applies functions  $f$  and  $g$  respectively to the elements of a pair  $(x, y)$  is denoted by  $(f \times g)$ . Thus,  $(f \times g)(x, y) = (f x, g y)$ . A function that applies functions  $f$  and  $g$  separately to an element and returns a pair of the results is denoted by  $(f \Delta g)$ . Thus,  $(f \Delta g) a = (f a, g a)$ . A projection function  $\pi_1$  extracts the first element of a pair. Thus,  $\pi_1(x, y) = x$ . These can be extended to the case of arbitrary number of elements.

Note that we use functional style notations only for parallel algorithm development; in fact we use the ordinary programming language C++ for practical coding.

## 2.1. TWO-DIMENSIONAL ARRAYS IN ABIDE-TREES

To represent two-dimensional arrays, we define the following abide-trees, which are built up by three constructors  $|\cdot|$  (singleton),  $\ominus$  (above) and  $\phi$  (beside) <sup>(7)</sup>.

```
data AbideTree  $\alpha$  =  $|\cdot|$   $\alpha$ 
                | (AbideTree  $\alpha$ )  $\ominus$  (AbideTree  $\alpha$ )
                | (AbideTree  $\alpha$ )  $\phi$  (AbideTree  $\alpha$ )
```

Here,  $|\cdot| a$ , or abbreviated as  $|a|$ , means a singleton array of  $a$ , i.e. a two-dimensional array with a single element  $a$ . We define the function `the` to extract the element from a singleton array, i.e. `the |a| = a`. For two-dimensional arrays  $x$  and  $y$  of the same width,  $x \ominus y$  means that  $x$  is located above  $y$ . Similarly, for two-dimensional arrays  $x$  and  $y$  of the same height,  $x \phi y$  means that  $x$  is located on the left of  $y$ . Moreover,  $\ominus$

and  $\phi$  are associative binary operators and satisfy the following *abide* (a coined term from above and beside) property.

DEFINITION 2.1 (Abide Property). *Two binary operators  $\oplus$  and  $\otimes$  are said to satisfy the abide property or to be abiding, if the following equation is satisfied:*

$$(x \otimes u) \oplus (y \otimes v) = (x \oplus y) \otimes (u \oplus v) .$$

In the rest of the paper, we will assume no inconsistency in height or width when  $\phi$  and  $\vartheta$  are used.

Note that one two-dimensional array may be represented by more than one abide-trees, but these abide-trees are equivalent because of the abide property of  $\vartheta$  and  $\phi$ . For example, we can express the  $2 \times 2$  two-dimensional array

$$\begin{pmatrix} 1 & 2 \\ 3 & 4 \end{pmatrix}$$

by the following two equivalent abide-trees.

$$\begin{aligned} &(|1| \phi |2|) \vartheta (|3| \phi |4|) \\ &(|1| \vartheta |3|) \phi (|2| \vartheta |4|) \end{aligned}$$

This is in sharp contrast to the quadtree representation of matrices<sup>(18)</sup>, which does not allow such freedom. This freedom is important in our framework. In our framework, a program construction consists of two phases. First, we make a general program that is architecture independent. Then, we derive a good program that may be architecture dependent or may have restricted order of access. In this case, restric-

tive representation prevents us from describing and deriving efficient programs. For example, a list of lists representation, which restricts the access order to outer dimension to inner dimension, does not allow us to describe and to derive efficient blocked algorithms. Thus, we start with a program using the abide-tree representation that does not impose restrictions on the access order, then we transform it to a good program that may have the restricted order of accesses. Moreover, this freedom allows easy re-balancing of the tree in computations such as divide-and-conquer computations on abide-trees.

## 2.2. ABIDE-TREE HOMOMORPHISM

From the theory of constructive algorithmics <sup>(9)</sup>, it follows that each constructively built-up data structure (i.e., algebraic data structure) is equipped with a powerful computation pattern called homomorphism.

DEFINITION 2.2 ((Abide-tree) Homomorphism).

A function  $h$  is said to be abide-tree homomorphism, if it is defined as follows for a function  $f$  and some binary operators  $\oplus, \otimes$ .

$$\begin{aligned} h \ |a| &= f \ a \\ h \ (x \ominus y) &= h \ x \oplus \ h \ y \\ h \ (x \ \phi \ y) &= h \ x \ \otimes \ h \ y \end{aligned}$$

For notational convenience, we write  $(|f, \oplus, \otimes|)$  to denote  $h$ . When it is clear from the context, we just call  $(|f, \oplus, \otimes|)$  homomorphism.

Intuitively, a homomorphism  $(|f, \oplus, \otimes|)$  is a function to replace the constructors  $|\cdot|$ ,  $\ominus$  and  $\phi$  in an input abide-tree by  $f$ ,  $\oplus$  and  $\otimes$  re-

spectively. We will see in Section 3 that many algorithms on two-dimensional arrays can be concisely specified by homomorphisms.

Note that  $\oplus$  and  $\otimes$  in  $(f, \oplus, \otimes)$  should be associative and satisfy the abide property, inheriting the properties of  $\oplus$  and  $\otimes$ .

Homomorphism enjoys many nice transformation rules, among which the following fusion rule is of particular importance. The fusion rule gives us a way to create a new homomorphism from composition of a function and a homomorphism. As will be seen in Section 4, it plays a key role in derivation of efficient parallel programs on abide-trees.

**THEOREM 2.3 (Fusion).** Let  $h$  and  $(f, \oplus, \otimes)$  be given. If there exist  $\odot$  and  $\ominus$  such that for any  $x$  and  $y$ ,

$$\begin{cases} h(x \oplus y) = h x \odot h y \\ h(x \otimes y) = h x \ominus h y \end{cases}$$

hold, then

$$h \circ (f, \oplus, \otimes) = (h \circ f, \odot, \ominus) .$$

*Proof.* The theorem is proved by induction on the structure of abide-trees. See the technical report <sup>(16)</sup> for details.  $\square$

Because of the flexibility of the abide-tree representation, a homomorphism  $(f, \oplus, \otimes)$  can be implemented *efficiently* in parallel, which will be shown in Section 5. Therefore, we will design our parallel skeletons based on the homomorphisms.

### 3. Parallel Skeletons and Algorithms on Two-Dimensional Arrays

In this section, we design a set of basic parallel skeletons for manipulating two-dimensional arrays, and show composition of the skeletons is powerful enough to describe useful parallel algorithms. Since homomorphisms are suitable for manipulating and developing parallel programs, which has been argued in the previous section, we will use the homomorphisms for the basis of the design of our parallel skeletons. The power of composition of the skeletons will be shown by describing nontrivial problems such as matrix multiplication and QR decomposition. One important fact here is that any homomorphism can be written as a composition of two basic skeletons. This fact is motivating the systematic derivation of parallel programs in Section 4.

#### 3.1. DATA PARALLEL SKELETONS

We define four primitive functions `map`, `reduce`, `zipwith` and `scan` on the data type *AbideTree*. In the theory of Constructive Algorithmics (7,9,38), these functions are known to be the most fundamental computation components for manipulating algebraic data structures and for being glued together to express complicated computations. We call them *parallel skeletons* because they have potential parallelism and can be implemented efficiently in parallel (see Section 5.) Intuitive definitions of the skeletons are shown in Figure 1.

### Map and Reduce

The skeletons `map` and `reduce` are two special cases of homomorphism.

The skeleton `map` applies a function  $f$  to each element of a two-dimensional array while keeping the structure, and is defined by

$$\begin{aligned} \text{map } f \mid a \mid &= \mid f a \mid \\ \text{map } f (x \ominus y) &= (\text{map } f x) \ominus (\text{map } f y) \\ \text{map } f (x \oplus y) &= (\text{map } f x) \oplus (\text{map } f y) , \end{aligned}$$

that is,  $\text{map } f = (\mid \cdot \mid \circ f, \ominus, \oplus)$ . For example, a scalar product of an array is performed with `map` as follows.

$$\text{map } (2 \times) \begin{pmatrix} 6 & 2 & 1 \\ 4 & 3 & 5 \end{pmatrix} = \begin{pmatrix} 12 & 4 & 2 \\ 8 & 6 & 10 \end{pmatrix}$$

The skeleton `reduce` collapses a two-dimensional array to a value using two abiding binary operators  $\oplus$ ,  $\otimes$ , and is defined by

$$\begin{aligned} \text{reduce}(\oplus, \otimes) \mid a \mid &= a \\ \text{reduce}(\oplus, \otimes) (x \ominus y) &= (\text{reduce}(\oplus, \otimes) x) \oplus (\text{reduce}(\oplus, \otimes) y) \\ \text{reduce}(\oplus, \otimes) (x \oplus y) &= (\text{reduce}(\oplus, \otimes) x) \otimes (\text{reduce}(\oplus, \otimes) y) , \end{aligned}$$

that is,  $\text{reduce}(\oplus, \otimes) = (\text{id}, \oplus, \otimes)$ . For example, a summation of an array is calculated with `reduce` as follows.

$$\text{reduce}(+, +) \begin{pmatrix} 6 & 2 & 1 \\ 4 & 3 & 5 \end{pmatrix} = 21$$

Interestingly, any homomorphism can be written as a composition of `map` and `reduce`.

LEMMA 3.1 (Homomorphism). A homomorphism  $(f, \oplus, \otimes)$  can be written as a composition of `map` and `reduce`:

$$(f, \oplus, \otimes) = \text{reduce}(\oplus, \otimes) \circ \text{map } f .$$

*Proof.* The lemma is proved by induction on the structure of abidetrees.  $\square$

This lemma implies that if we have efficient parallel implementations for `reduce` and `map`, we get an efficient implementation for homomorphism. Since it is shown in the Section 5 that parallel skeletons have efficient parallel implementations, a homomorphism has an efficient parallel implementations. This fact is motivating the systematic derivation of parallel programs in Section 4.

### *Zipwith*

The two skeletons defined above are primitive skeletons. We define other skeletons that are extensions of these primitive skeletons. The skeleton `zipwith`, an extension of `map`, takes two two-dimensional arrays of the same shape, applies a function  $f$  to corresponding elements of the arrays and returns a new array of the same shape.

$$\begin{aligned} \text{zipwith } f \ |a| \ |b| &= |f \ a \ b| \\ \text{zipwith } f \ (x \oplus y) \ (u \oplus v) &= (\text{zipwith } f \ x \ u) \oplus (\text{zipwith } f \ y \ v) \\ \text{zipwith } f \ (x \ \phi \ y) \ (u \ \phi \ v) &= (\text{zipwith } f \ x \ u) \ \phi \ (\text{zipwith } f \ y \ v) \end{aligned}$$

Note that in the above definition two-dimensional arrays that are the arguments of the function should be divided in the way that the sizes of  $x$  and  $u$  are the same. For example, an addition of two arrays is

calculated with `reduce` as follows.

$$\text{zipwith}(+) \begin{pmatrix} 6 & 2 & 1 \\ 4 & 3 & 5 \end{pmatrix} \begin{pmatrix} 0 & 8 & 2 \\ 9 & 1 & 7 \end{pmatrix} = \begin{pmatrix} 6 & 10 & 3 \\ 13 & 4 & 12 \end{pmatrix}$$

Function `zip` is a specialization of `zipwith`, making a two-dimensional array of pairs of corresponding elements.

$$\text{zip}(u, v) = \text{zipwith}(\lambda xy. (x, y)) u v$$

We may define similar `zip` and `zipwith` for the case when the number of input arrays is three or more, and those that take  $k$  arrays are denoted by `zipk` and `zipwithk`. Also, we define `unzip` to be the inverse of `zip`.

Composing these skeletons defined above, we can describe many useful functions.

$$\begin{aligned}
id &= \text{reduce}(\ominus, \phi) \circ \text{map } |\cdot| \\
tr &= \text{reduce}(\phi, \ominus) \circ \text{map } |\cdot| \\
rev &= \text{reduce}(\tilde{\ominus}, \tilde{\phi}) \circ \text{map } |\cdot| \\
flatten &= \text{reduce}(\ominus, \phi) \\
height &= \text{reduce}(+, \ll) \circ \text{map } (\lambda x. 1) \\
width &= \text{reduce}(\ll, +) \circ \text{map } (\lambda x. 1) \\
cols &= \text{reduce}(\text{zipwith}(\ominus), \phi) \circ \text{map } ||\cdot|| \\
rows &= \text{reduce}(\ominus, \text{zipwith}(\phi)) \circ \text{map } ||\cdot|| \\
\text{reduce}_c(\oplus) &= \text{map}(\text{reduce}(\oplus, \ll)) \circ cols \\
\text{reduce}_r(\otimes) &= \text{map}(\text{reduce}(\ll, \otimes)) \circ rows \\
\text{map}_c f &= \text{reduce}(\ll, \phi) \circ \text{map } f \circ cols \\
\text{map}_r f &= \text{reduce}(\ominus, \ll) \circ \text{map } f \circ rows \\
add &= \text{zipwith}(+) \\
sub &= \text{zipwith}(-)
\end{aligned}$$

Here,  $||\cdot||$  is abbreviation of  $|\cdot| \circ |\cdot|$ . The function *id* is the identity function of *AbideTree*, and *tr* is the matrix-transposing function. The function *rev* takes a two-dimensional array and returns the array reversed in the vertical and the horizontal direction, and *flatten* flattens a nested *AbideTree*. The functions *height* and *width* return the number of rows and columns respectively, and *cols* and *rows* return an array of which elements are columns and rows of the array of the argument respectively. The functions  $\text{reduce}_c$  and  $\text{reduce}_r$  that are specializations of *reduce* reduce a two-dimensional array in each column and row direction respectively and return a row-vector (an array of which height is one)

and a column-vector (an array of which width is one). The functions  $\text{map}_c$  and  $\text{map}_r$  that are specializations of  $\text{map}$  apply a function to each column and row respectively (i.e. the function of the argument takes column-vector or row-vector). The functions  $\text{add}$  and  $\text{sub}$  denote matrix addition and subtraction respectively.

### *Scan*

The skeleton  $\text{scan}$ , an extension of  $\text{reduce}$ , holds all values generated in reducing a two-dimensional array by  $\text{reduce}$ .

$$\begin{aligned}\text{scan}(\oplus, \otimes) |a| &= |a| \\ \text{scan}(\oplus, \otimes)(x \ominus y) &= (\text{scan}(\oplus, \otimes) x) \oplus' (\text{scan}(\oplus, \otimes) y) \\ \text{scan}(\oplus, \otimes)(x \oplus y) &= (\text{scan}(\oplus, \otimes) x) \otimes' (\text{scan}(\oplus, \otimes) y)\end{aligned}$$

Here operators  $\oplus'$  and  $\otimes'$  are defined as follows.

$$\begin{aligned}sx \oplus' sy &= sx \ominus sy' \\ &\textbf{where } sy' = \text{map}_r(\text{zipwith}(\oplus)(\text{bottom } sx)) sy \\ sx \otimes' sy &= sx \oplus sy' \\ &\textbf{where } sy' = \text{map}_c(\text{zipwith}(\otimes)(\text{last } sx)) sy \\ \text{bottom} &= \text{reduce}(\gg, \oplus) \circ \text{map } |\cdot| \\ \text{last} &= \text{reduce}(\ominus, \gg) \circ \text{map } |\cdot|\end{aligned}$$

For example, a prefix-um (an upper-left prefix-sum) of an array is calculated with  $\text{scan}$  as follows.

$$\text{scan}(+, +) \begin{pmatrix} 6 & 2 & 1 \\ 4 & 3 & 5 \end{pmatrix} = \begin{pmatrix} 6 & 8 & 9 \\ 10 & 15 & 21 \end{pmatrix}$$

It should be noted that `reduce` can be expressed by `reducec` and `reducer` when two binary operators  $\oplus$  and  $\otimes$  are abiding.

$$\begin{aligned} \text{reduce}(\oplus, \otimes) &= \text{the} \circ \text{reduce}_c(\oplus) \circ \text{reduce}_r(\otimes) \\ \text{reduce}(\oplus, \otimes) &= \text{the} \circ \text{reduce}_r(\otimes) \circ \text{reduce}_c(\oplus) \end{aligned} \quad (1)$$

Like `reduce`, we may define `scan↓` and `scan→` that are specialization of `scan` and scan a two-dimensional array in column and row direction respectively:

$$\begin{aligned} \text{scan}_{\downarrow}(\oplus) &= \text{scan}(\oplus, \gg) \\ \text{scan}_{\rightarrow}(\otimes) &= \text{scan}(\gg, \otimes) ; \end{aligned}$$

`scan` can be expressed by `scan↓` and `scan→` when two binary operators  $\oplus$  and  $\otimes$  are abiding.

$$\begin{aligned} \text{scan}(\oplus, \otimes) &= \text{scan}_{\downarrow}(\oplus) \circ \text{scan}_{\rightarrow}(\otimes) \\ \text{scan}(\oplus, \otimes) &= \text{scan}_{\rightarrow}(\otimes) \circ \text{scan}_{\downarrow}(\oplus) \end{aligned} \quad (2)$$

Using the skeleton `scan`, we can define `scanr` that executes `scan` reversely (i.e. from bottom to top, from right to left), `allredr` and `allredc` that broadcast the results in each row and column after `reducer` and `reducec` respectively. These functions are used in later section.

$$\begin{aligned} \text{scanr}(\oplus, \otimes) &= \text{rev} \circ \text{scan}(\tilde{\oplus}, \tilde{\otimes}) \circ \text{rev} \\ \text{allred}_c(\oplus) &= \text{scanr}(\gg, \ll) \circ \text{scan}(\oplus, \gg) \\ \text{allred}_r(\otimes) &= \text{scanr}(\ll, \gg) \circ \text{scan}(\gg, \otimes) \end{aligned}$$

### 3.2. DATA COMMUNICATION SKELETONS

We show how to define data communication skeletons **dist**, **gather**,  $\text{rot}_r$  and  $\text{rot}_c$  that abstract distribution, collection and rearrangement of a two-dimensional array among processors. The idea is to use nested two-dimensional arrays to represent distributed two-dimensional arrays.

The skeleton **dist** abstracts distribution of a two-dimensional array to processors, and is defined as

$$\begin{aligned} \mathbf{dist} \, p \, q \, x &= (\mathit{flatten} \circ \mathbf{map}(\mathit{grp}_c \, n) \circ \mathit{grp}_r \, m) \, x \\ &\mathbf{where} \, m = \lceil \mathit{height} \, x / p \rceil, \, n = \lceil \mathit{width} \, x / q \rceil \end{aligned}$$

where  $\mathit{grp}_r$  is defined as follows and  $\mathit{grp}_c$  is defined similarly.

$$\begin{aligned} \mathit{grp}_r \, k \, (x \oplus y) &= |x| \oplus (\mathit{grp}_r \, k \, y) \quad \text{if } \mathit{height} \, x = k \\ \mathit{grp}_r \, k \, x &= |x| \quad \quad \quad \text{if } \mathit{height} \, x < k \end{aligned}$$

The distribution  $\mathbf{dist} \, p \, q \, x$  means that the two-dimensional array  $x$  will be divided into  $p \times q$  subarrays (i.e.  $x$  is divided into  $p$  subarrays in vertical direction, then each subarray is divided into  $q$  subarrays in horizontal direction), and each subarray is distributed to a processor.

The skeleton **gather**, the inverse operator of **dist**, abstracts gathering of two-dimensional arrays distributed to the processors into a two-dimensional array on the root processor.

$$\mathbf{gather} = \mathbf{reduce}(\oplus, \phi)$$

Although definitions of these skeletons may seem complicated, actual operations are simple as illustrated in Figure 2. What is significant here is that these skeletons satisfy the relation of  $id = \mathbf{gather} \circ \mathbf{dist} \ p \ q$ .

The rotation skeleton  $\mathbf{rot}_r$  that takes a function  $f$  and rotates the  $i$ -th row (the index begins from 0) by  $f \ i$ , is defined as follows:

$$\mathbf{rot}_r \ f = \mathbf{flatten} \circ \mathbf{map} \ \mathbf{shift}_r \circ \mathbf{addidx}_r \circ \mathbf{rows}$$

**where**

$$\mathbf{addidx}_r \ u = \mathbf{zip}(\mathbf{map} \ f \ (\mathbf{idx}_r \ u), u)$$

$$\mathbf{idx}_r = \mathbf{map}(-1) \circ \mathbf{scan}_\downarrow(+)$$

$$\circ \mathbf{map}(\lambda x. 1) \ ;$$

here  $\mathbf{shift}_r$  is defined under the condition  $i > 0$ .

$$\mathbf{shift}_r(0, x) = x$$

$$\mathbf{shift}_r(i, x \ \phi \ y) = y \ \phi \ x \ \text{if } \mathit{width} \ y = i$$

$$\mathbf{shift}_r(-i, x \ \phi \ y) = y \ \phi \ x \ \text{if } \mathit{width} \ x = i$$

Similarly, we can define the skeleton  $\mathbf{rot}_c$  that takes a function  $f$  and rotates the  $i$ -th column by  $f \ i$ . An image of the above communication skeletons is depicted in Figure 2. In the figure, since the rotation skeleton  $\mathbf{rot}_r$  takes a negation function, the 0-th row does not rotate (rotates by 0), the first row rotates to the left by 1 (to the right by  $-1$ ) and the second row rotates to the left by 2 (to the right by  $-2$ ).

### 3.3. MATRIX MULTIPLICATION

As an involved example, we describe two known parallel algorithms for matrix multiplication, which is a primitive operation of matrices, with the above defined parallel skeletons on two-dimensional arrays.

The first description is Cannon's Algorithm <sup>(22)</sup>:

$$mm_C = \text{gather} \circ (\text{map } thd) \circ (\text{iter } p \text{ step}) \circ \text{arrange} \circ \text{distribute} \circ \text{init}$$

**where**

$$\text{init } (A, B) = (A, B, \text{map}(\lambda x. 0) A)$$

$$\text{distribute} = (\text{dist } p p \times \text{dist } p p \times \text{dist } p p)$$

$$\text{arrange} = \text{zip}_3 \circ (\text{rot}_r \text{ neg} \times \text{rot}_c \text{ neg} \times \text{id})$$

$$\text{step} = \text{rearrange} \circ \text{unzip}_3 \circ \text{map } lmm$$

$$\text{rearrange} = \text{zip}_3 \circ (\text{rot}_r(\lambda x. 1) \times \text{rot}_c(\lambda x. 1) \times \text{id})$$

$$\text{neg } x = -x$$

$$thd (x, y, z) = z$$

where  $p$  is a natural number indicating the number of divisions of matrices in column and row direction, and  $lmm$  is a function that executes locally matrix multiplication on matrices on each processor, i.e.  $lmm (A, B, C) = (A, B, C + A \times B)$ . The function  $iter$  is defined as follows.

$$\text{iter } k \text{ f } x = x \quad \text{if } k = 0$$

$$\text{iter } k \text{ f } x = \text{iter } (k - 1) \text{ f } (f x) \quad \text{if } k > 0$$

Explicit distribution of matrices by the data communication skeletons makes this description looking complicated. However, it should be noted that even non-intuitive complicated Cannon's Algorithm can be described by composition of the skeletons.

The second description is an intuitively understandable description using only data parallel skeletons (an element of the resulting matrix is an inner product of a row vector and a column vector of the input matrices.) This description describes just a definition of matrix multi-

plication. Although users do not need to consider parallelism at all, this program can be executed in parallel due to parallelism of each skeleton.

$$mm = \text{zipwith}_P \text{ iprod} \circ (\text{allrows} \times \text{allcols})$$

**where**

$$\text{allrows} = \text{allred}_r(\phi) \circ \text{map} |\cdot|$$

$$\text{allcols} = \text{allred}_c(\ominus) \circ \text{map} |\cdot|$$

$$\text{iprod} = (\text{reduce}(+, +) \circ) \circ \text{zipwith}(\times) \circ \text{tr}$$

$$\text{zipwith}_P(\otimes)(x, y) = \text{zipwith}(\otimes) x y$$

Although this definition seems to use  $O(n^3)$  memory space for  $n \times n$  matrices due to duplications of rows and columns with  $\text{allred}_r$  and  $\text{allred}_c$ , we can execute this multiplication using  $O(n^2)$  memory space. This is because we can use references instead of duplications in the implementation of  $\text{allred}_r$  and  $\text{allred}_c$  due to the properties of the operators in their definitions. This kind of optimization is currently done by hand. However, we think it will be automatically done by compilers when it takes the properties of the operators into account.

### 3.4. QR FACTORIZATION

As the final nontrivial example, we show descriptions of two parallel algorithms for QR factorization <sup>(15)</sup>. We will not explain the details, but we hope to show that these algorithms can be dealt with in our framework.

We give the recursive description of a QR factorization algorithm based on Householder transform. This function returns  $Q$  and  $R$  which satisfy  $A = QR$  where  $A$  is a matrix of  $m \times n$ ,  $Q$  an orthogonal matrix

of  $m \times m$  and  $R$  an upper-triangular matrix of  $m \times n$ .

$$\begin{aligned}
& qr ((A_{11} \oplus A_{21}) \oplus (A_{12} \oplus A_{22})) \\
& = \mathbf{let} (Q_1, R_{11} \oplus 0) = qr (A_{11} \oplus A_{21}) \\
& \quad (R_{12} \oplus \hat{A}_{22}) = mm (tr Q_1) (A_{12} \oplus A_{22}) \\
& \quad (\hat{Q}_2, R_{22}) = qr \hat{A}_{22} \\
& \quad Q = mm Q_1 ((I \oplus 0) \oplus (0 \oplus \hat{Q}_2)) \\
& \quad \mathbf{in} (Q, (R_{11} \oplus R_{12}) \oplus (0 \oplus R_{22})) \\
& qr (|a| \oplus x) = hh (|a| \oplus x) \\
& hh v \quad = \mathbf{let} v' = add v e \\
& \quad a = \sqrt{\text{reduce}(+, +) (\text{zipwith}(\times) v' v')} \\
& \quad u = \text{map} (/a) v' \\
& \quad Q = \text{sub } I (\text{map} (\times 2) (mm u (tr u))) \\
& \quad \mathbf{in} (Q, e)
\end{aligned}$$

Here  $e$  is a vector (a matrix of which width is 1) whose first element is 1 and the other elements are 0, and  $I$  and  $0$  represent an identity matrix and a zero matrix of suitable size respectively.

Furthermore, we give the recursive description of QR factorization algorithm on quadtree <sup>(18)</sup>; transforming algorithms on quadtrees to those on abide-trees is always possible because abide-trees is more flexible than quadtrees. This function  $qr_q$  is mutual recursively defined with an extra function  $e$ , and returns  $Q$  and  $R$  that satisfy  $A = QR$  where  $A$  is a matrix of  $n \times n$  ( $n = 2^k$  for a natural number  $k$ ),  $Q$  is an orthogonal matrix of  $n \times n$  and  $R$  is an upper-triangular matrix of

$n \times n$ .

$$\begin{aligned}
qr_q |a| &= (|1|, |a|) \\
qr_q ((A_{11} \oplus A_{21}) \oplus (A_{12} \oplus A_{22})) \\
&= \mathbf{let} (Q_1, R_1) = qr_q A_{11} \\
&\quad (Q_2, R_2) = qr_q A_{21} \\
&\quad Q_{12} = (Q_1 \oplus 0) \oplus (0 \oplus Q_2) \\
&\quad (Q_3, R_3) = e (R_1, R_2) \\
&\quad Q_4 = mm Q_{12} Q_3 \\
&\quad (U_n \oplus U_s) = mm (tr Q_4) (A_{12} \oplus A_{22}) \\
&\quad (Q_6, R_6) = qr_q U_s \\
&\quad Q = mm Q_4 ((I \oplus 0) \oplus (0 \oplus Q_6)) \\
&\quad R = (R_3 \oplus U_n) \oplus (0 \oplus R_6) \\
&\mathbf{in} (Q, R)
\end{aligned}$$

Note that  $A_{ij}$  ( $i, j \in \{1, 2\}$ ) have the same shape. A definition of the involved extra function  $e$  is as follows.

$$\begin{aligned}
e(N, O) &= (I, N) \\
e(|n|, |s|) &= \mathbf{let} \ Q = g(n, s) \\
&\quad (N, O) = mm(tr \ Q) (|n| \oplus |s|) \\
&\quad \mathbf{in} \ (Q, N) \\
e((N_{11} \oplus N_{21}) \oplus (N_{12} \oplus N_{22}), (S_{11} \oplus S_{21}) \oplus (S_{12} \oplus S_{22})) \\
&= \mathbf{let} \\
&\quad ((Q_1^{11} \oplus Q_1^{21}) \oplus (Q_1^{12} \oplus Q_1^{22}), N_1) = e(N_{11}, S_{11}) \\
&\quad ((Q_2^{11} \oplus Q_2^{21}) \oplus (Q_2^{12} \oplus Q_2^{22}), N_2) = e(N_{22}, S_{22}) \\
Q_{12} &= (Q_1^{11} \oplus O \oplus Q_1^{12} \oplus O) \oplus (O \oplus Q_2^{11} \oplus O \oplus Q_2^{12}) \\
&\quad \oplus (Q_1^{21} \oplus O \oplus Q_1^{22} \oplus O) \oplus (O \oplus Q_2^{21} \oplus O \oplus Q_2^{22}) \\
Q_1 &= (Q_1^{11} \oplus Q_1^{21}) \oplus (Q_1^{12} \oplus Q_1^{22}) \\
(U_n \oplus U_s) &= mm(tr \ Q_1) (N_{12} \oplus S_{12}) \\
(Q_4, R_4) &= qr_q \ U_s \\
Q'_4 &= (I \oplus O \oplus O \oplus O) \oplus (O \oplus I \oplus O \oplus O) \\
&\quad \oplus (O \oplus O \oplus Q_4 \oplus O) \oplus (O \oplus O \oplus O \oplus I) \\
Q_5 &= mm \ Q_{12} \ Q'_4 \\
((Q_6^{11} \oplus Q_6^{21}) \oplus (Q_6^{12} \oplus Q_6^{22}), N_6) &= e(N_2, R_4) \\
Q'_6 &= (I \oplus O \oplus O \oplus O) \oplus (O \oplus Q_6^{11} \oplus Q_6^{12} \oplus O) \\
&\quad \oplus (O \oplus Q_6^{21} \oplus Q_6^{22} \oplus O) \oplus (O \oplus O \oplus O \oplus I) \\
&\mathbf{in} \ (mm \ Q_5 \ Q'_6, (N_1 \oplus U_n) \oplus (O \oplus N_6)) \\
g(a, b) &= (|c| \oplus |s|) \oplus (|-s| \oplus |c|) \\
&\quad \mathbf{where} \ c = \frac{a}{\sqrt{a^2 + b^2}}, \ s = \frac{-b}{\sqrt{a^2 + b^2}}
\end{aligned}$$

Note that  $N_{ij}$  and  $S_{ij}$  ( $i, j \in \{1, 2\}$ ) have the same shape and  $Q_k^{ij}$  ( $i, j, k \in \{1, 2\}$ ) have the same shape.

We can efficiently parallelize some parts of these complicated recursive functions, such as matrix multiplication and independent calculations like  $(Q_1, R_1) = qr_q A_{11}$  and  $(Q_2, R_2) = qr_q A_{21}$ . These independent calculations are explicitly parallelized by describing them with the `map` skeleton as follows.

$$\left\{ \begin{array}{l} (Q_1, R_1) = qr_q A_{11} \\ (Q_2, R_2) = qr_q A_{21} \end{array} \right\}$$

↓

$$|(Q_1, R_1)| \oplus |(Q_2, R_2)| = \text{map } \textit{apply} (|(qr_q, A_{11})| \oplus |(qr_q, A_{21})|)$$

Here, the function *apply* is defined as  $\textit{apply} (f, x) = f x$ . It is, however, still an open problem whether the complicated recursive functions can be parallelized with our defined skeletons, although we can introduce other skeletons such as divide-and-conquer skeleton to parallelize them. This is our future work.

#### 4. Developing Efficient Parallel Programs

It has been shown so far that compositions of recursive functions on abide-trees including homomorphisms and skeletons provide us with a powerful mechanism to describe parallel algorithms on two-dimensional arrays, where parallelism in the original parallel algorithms can be well

captured. In this section, we move on from issues of parallelism to the issues of efficiency. We will illustrate a strategy to guide programmers to develop *efficient* parallel algorithms systematically through program transformation.

Recall homomorphisms have efficient parallel implementation as composition of our parallel skeletons. Thus, a goal of this derivation may be to write a program by a homomorphism. However, not all functions can be specified by a single homomorphism. Therefore, we first introduce a more powerful tool called almost-homomorphism <sup>(12)</sup>. Then, we demonstrate the strategy with an example.

#### 4.1. ALMOST-HOMOMORPHISM

Not all functions can be specified by a single homomorphism, but we can always tuple these functions with some extra functions so that the tupled functions can be specified by a homomorphism. An *almost homomorphism* is a composition of a projection function and a homomorphism. Since projection functions are simple, almost homomorphisms are suitable for parallel computation as homomorphisms are.

In fact, every function can be represented in terms of an almost homomorphism at the cost of redundant computation. Let  $k$  be a nonhomomorphic function, and consider a new function  $g$  such that

$g x = (k x, x)$ . The tupled function  $g$  is a homomorphism.

$$g |a| = (k |a|, |a|)$$

$$g (x \oplus y) = g x \oplus g y$$

$$\textbf{where } (k_1, x_1) \oplus (k_2, x_2) = (k (x_1 \oplus x_2), x_1 \oplus x_2)$$

$$g (x \otimes y) = g x \otimes g y$$

$$\textbf{where } (k_1, x_1) \otimes (k_2, x_2) = (k (x_1 \otimes x_2), x_1 \otimes x_2)$$

Then,  $k$  is written as an almost homomorphism:

$$k = \pi_1 \circ g = \pi_1 \circ (g \circ |\cdot|, \oplus, \otimes) .$$

However, the definition above is not efficient because binary operators  $\oplus$  and  $\otimes$  do not use the previously computed values  $k_1$  and  $k_2$ . In order to derive a good almost homomorphism, we should carefully define a suitable tupled function, making full use of previously computed values. We will see this in our parallel program development in the next section. However, to determine the class of problems that have good (efficient) almost-homomorphic implementation is an open problem.

#### 4.2. A STRATEGY FOR DERIVING EFFICIENT PARALLEL PROGRAMS

Our strategy for deriving efficient parallel programs on two-dimensional arrays consists of the following four steps, extending the result of lists <sup>(24)</sup>. The goal of the strategy is to write a given program by an efficient almost-homomorphism that has an efficient implementation in parallel.

Step 1. Define the target program  $p$  as a composition of  $p_1, \dots, p_n$  that are already defined, i.e.  $p = p_n \circ \dots \circ p_1$ . Each of  $p_1, \dots, p_n$  may

be defined as a composition of small functions or a recursive function (see Section 3.3 and Section 3.4).

Step 2. Derive an almost homomorphism (Section 4.1) from the recursive definition of  $p_1$ .

Step 3. Fuse  $p_2$  into the derived almost homomorphism to obtain a new almost homomorphism for  $p_2 \circ p_1$ , and repeat this derivation until  $p_n$  is fused.

Step 4. Let  $\pi_1 \circ (f, \oplus, \otimes)$  be the resulting almost homomorphism for  $p_n \circ \dots \circ p_1$  obtained at Step 3. For the functions inside the homomorphism, namely  $f$ ,  $\oplus$  and  $\otimes$ , try to repeat Steps 2 and 3 to find efficient parallel implementations for them.

In the following, we explain this strategy through a derivation of an efficient program for the maximum rectangle sum problem: compute the maximum of sums of all the rectangle data areas in a two-dimensional data. This problem was originated by Bentley <sup>(2,3)</sup> and improved by Takaoka <sup>(41)</sup>. The solution can be used in a sort of data mining and pattern matching of two dimensional data. For example, for the following two-dimensional data

$$\begin{pmatrix} 3 & -1 & 4 & -1 & -5 \\ 1 & -4 & -1 & \mathbf{5} & -3 \\ -4 & 1 & \mathbf{5} & \mathbf{3} & 1 \end{pmatrix}$$

the result should be 15, which denotes the maximum sum contributed by the sub-rectangular area with bolded numbers above. To appreciate

difficulty of this problem, we ask the reader to pause for a while to think of how to solve it.

We will omit some portions of the description in the following derivation for the readability. Please see the technical report <sup>(16)</sup> for the full description.

#### STEP 1. DEFINING A CLEAR PARALLEL PROGRAM

A clear and straightforward solution to the maximum rectangle sum problem is as follows: enumerating all possible rectangles, then computing sums for all rectangles, and finally returning the maximum value as the result.

$$mrs = max \circ \text{map } max \circ \text{map } (\text{map } sum) \circ rects$$

**where**

$$max = \text{reduce}(\uparrow, \uparrow)$$

$$sum = \text{reduce}(+, +)$$

Here *rects* is a function that takes a two-dimensional array and returns all possible rectangles of the array. The returned value of *rects* is an array of arrays of arrays, and the  $(k, l)$ -element of the  $(i, j)$ -element of the resulting array is a sub-rectangle having rows from the  $i$ -th to the  $j$ -th and columns from the  $k$ -th to the  $l$ -th of the original array. An example of *rects* is shown below. Note that we think that the special value is contained in the blank portion of the above-mentioned array, and we write the blank of arbitrary size by *NIL* for brevity. Here, *NIL*

may be an array of which element is  $-\infty$  or an array of it.

$$\mathit{rects} \begin{pmatrix} 1 & 2 & 3 \\ 5 & 6 & 7 \end{pmatrix} = \left( \left( \begin{pmatrix} (1) \\ (2) \\ (3) \end{pmatrix} \begin{pmatrix} (1\ 2) \\ (2\ 3) \end{pmatrix} \begin{pmatrix} (1\ 2\ 3) \end{pmatrix} \right) \left( \begin{pmatrix} (1) \\ (5) \end{pmatrix} \begin{pmatrix} (1\ 2) \\ (5\ 6) \\ (2) \\ (6) \end{pmatrix} \begin{pmatrix} (1\ 2\ 3) \\ (5\ 6\ 7) \\ (2\ 3) \\ (6\ 7) \\ (3) \\ (7) \end{pmatrix} \right) \right)$$

The function *rects* is mutual recursively defined as follows.

$$\begin{aligned} \mathit{rects} \ |a| &= |||a||| \\ \mathit{rects} \ (x \ominus y) &= (\mathit{rects} \ x \ \phi \ \mathit{gemm}(\_, \mathit{zipwith}(\ominus)) \ (\mathit{bottoms} \ x) \ (\mathit{tops} \ y)) \\ &\quad \ominus (\mathit{NIL} \ \phi \ \mathit{rects} \ y) \\ \mathit{rects} \ (x \ \phi \ y) &= \mathit{zipwith}_4 \ f_s \ (\mathit{rects} \ x) \ (\mathit{rects} \ y) \ (\mathit{rights} \ x) \ (\mathit{lefts} \ y) \\ \mathbf{where} \ f_s \ s_1 \ s_2 \ r_1 \ l_2 &= (s_1 \ \phi \ \mathit{gemm}(\_, \ \phi) \ r_1 \ l_2) \ominus (\mathit{NIL} \ \phi \ s_2) \end{aligned}$$

where ‘\_’ indicates “don’t care” and generalized matrix multiplication *gemm* is defined as follows.

$$gemm(\oplus, \otimes) = g$$

**where**

$$g(X_1 \oplus X_2)(Y_1 \ominus Y_2) = zipwith(\oplus)(g X_1 Y_1)(g X_2 Y_2)$$

$$g(X_1 \ominus X_2)Y = (g X_1 Y) \ominus (g X_2 Y)$$

$$gX(Y_1 \oplus Y_2) = (g X Y_1) \oplus (g X Y_2)$$

$$g|a||b| = |a \otimes b|$$

Figure 3 shows recursive computation of *rects* for the  $\ominus$  case. Since indices  $i$  and  $j$  of the resulting array mean the top row and the bottom row of the generated rectangles within the argument array, the upper-left block ( $i \leq m_x$  and  $j \leq m_x$ ) of the resulting array contains rectangles included in the upper block of the argument, i.e.  $x$ . Thus, the upper-left block is *rects x*, and the lower-right block is *rects y* similarly (Figure 3-(a)). Each rectangle in the the upper-right block ( $i \leq m_x$  and  $j > m_x$ ) consists of a rectangle at the bottom of  $x$  and a rectangle at the top of  $y$  as shown in Figure 3-(b). Since there are all combinations of  $i$  and  $j$ , this block is computed by a general matrix multiplication of *bottoms x* and *tops y* as shown in Figure 3-(c). The computation of the  $\ominus$  case is similar.

Functions *bottoms*, *tops*, *rights* and *lefts* are similarly defined as mutual recursive functions shown in Figure 5, and their examples are shown in Figure 6. Each of these functions is the partial result of *rects* in the sense that it returns the rectangles that are restricted to include segments of some edges of the input as shown in Figure 8. For example, the *tops* returns the rectangles that include segments of the top edge

(the top-most row) of the input array. Similarly, *bottoms*, *rights* and *lefts* return the rectangles that include segments of the bottom edge, the right edge and the left edge respectively. The other functions returns the rectangles that include segments of two edges of the input. Usually, users do not need to understand well these functions because these functions often used are provided by experts.

Although this initial program is clear and has all its parallelism specified in terms of our parallel skeletons, it is inefficient in the sense that it needs to execute  $O(n^6)$  addition operations for the input of  $n \times n$  array. We will show how to develop a more efficient parallel program.

## STEP 2. DRIVING ALMOST HOMOMORPHISM

First, we propose a way of deriving almost homomorphism from mutual recursive definitions. For notational convenience, we define

$$\begin{aligned}\Delta_1^n f_i &= f_1 \Delta f_2 \Delta \cdots \Delta f_n \\ x(\Delta_1^n \oplus_i) y &= (x \oplus_1 y, x \oplus_2 y, \dots, x \oplus_n y) .\end{aligned}$$

Our main idea is based on the following theorem.

**THEOREM 4.1** (Tupling).

Let  $h_1, h_2, \dots, h_n$  be mutual recursively defined by

$$\begin{cases} h_i |a| &= f_i a , \\ h_i (x \oplus y) &= ((\Delta_1^n h_i) x) \oplus_i ((\Delta_1^n h_i) y) , \\ h_i (x \phi y) &= ((\Delta_1^n h_i) x) \otimes_i ((\Delta_1^n h_i) y) . \end{cases} \quad (3)$$

Then  $\Delta_1^n h_i$  is a homomorphism  $(\Delta_1^n f_i, \Delta_1^n \oplus_i, \Delta_1^n \otimes_i)$ .

*Proof.* The theorem is proven based on the definition of homomorphisms. See the technical report <sup>(16)</sup> for details.  $\square$

Theorem 4.1 says that if  $h_1$  is mutually defined with other functions (i.e.  $h_2, \dots, h_n$ ) which traverse over the same array in the specific form of Eq. (3), then tupling  $h_1, \dots, h_n$  will give a homomorphism. It follows that every  $h_i$  is an almost homomorphism. Thus, this theorem gives us a systematic way to execute Step 2 of the strategy.

We apply this theorem to derive an almost homomorphism for *rects*. The definition of *rects* and the extra functions are in the form of Eq. (3). Thus, we can obtain the following almost homomorphism by tupling these functions as follows (we show only operators corresponding to *rects* for readability.)

$$rects = \pi_1 \circ (\Delta_1^{11} f_i, \Delta_1^{11} \oplus_i, \Delta_1^{11} \otimes_i)$$

**where**

$$f_1 |a| = |||a|||$$

$$(s_1, t_1, b_1, r_1, l_1, tr_1, br_1, tl_1, bl_1, c_1, ro_1)$$

$$\oplus_1 (s_2, t_2, b_2, r_2, l_2, tr_2, br_2, tl_2, bl_2, c_2, ro_2)$$

$$= (s_1 \phi gemm(\_, zipwith(\oplus)) b_1 t_2) \oplus (NIL \phi s_2)$$

$$(s_1, t_1, b_1, r_1, l_1, tr_1, br_1, tl_1, bl_1, c_1, ro_1)$$

$$\otimes_1 (s_2, t_2, b_2, r_2, l_2, tr_2, br_2, tl_2, bl_2, c_2, ro_2)$$

$$= zipwith_4 f_s s_1 s_2 r_1 l_2$$

$$\mathbf{where} f_s s_1 s_2 r_1 l_2 = (s_1 \phi gemm(\_, \phi) r_1 l_2) \oplus (NIL \phi s_2)$$

These operators are straightforward rewriting of the definition of *rects*.

## STEP 3. FUSING WITH ALMOST HOMOMORPHISMS

We aim to derive an efficient almost homomorphism for *mrs*. To this end, we give the following theorem showing how to fuse a function with an almost homomorphism to get new another almost homomorphism.

**THEOREM 4.2 (Almost Fusion).**

Let  $h$  and  $(\Delta_1^n f_i, \Delta_1^n \oplus_i, \Delta_1^n \otimes_i)$  be given. If there exist  $\odot_i, \ominus_i$  ( $i = 1, \dots, n$ ) and  $H = h_1 \times h_2 \times \dots \times h_n$  ( $h_1 = h$ ) such that for any  $i$ ,  $x$  and  $y$ ,

$$h_i(x \oplus_i y) = H x \odot_i H y$$

$$h_i(x \otimes_i y) = H x \ominus_i H y$$

hold, then

$$h \circ (\pi_1 \circ (\Delta_1^n f_i, \Delta_1^n \oplus_i, \Delta_1^n \otimes_i)) = \pi_1 \circ (\Delta_1^n (h_i \circ f_i), \Delta_1^n \odot_i, \Delta_1^n \ominus_i) . \quad (4)$$

*Proof.* The theorem is proven by some calculation and Theorem 2.3. See the technical report <sup>(16)</sup> for details.  $\square$

Theorem 4.2 says that we can fuse a function with an almost homomorphism to get another almost homomorphism by finding  $h_2, \dots, h_n$  together with  $\odot_1, \dots, \odot_n, \ominus_1, \dots, \ominus_n$  that satisfy Eq. (4). Thus, this theorem gives us a systematic way to execute Step 3 of the strategy.

Returning to our example, we apply this theorem to *mrs*. The second function  $p_2$  of our example is  $\text{map}(\text{map } \text{sum})$ , so  $h_1 = \text{map}(\text{map } \text{sum})$ .

Then, we calculate  $h_1 (x \oplus_1 y)$  to find other functions and operators.

$$\begin{aligned}
& h_1 (x \oplus_1 y) \\
= & \quad \{ \text{Expand } x, y \text{ and } h_1 \} \\
& \text{map}(\text{map } \text{sum})((s_1 \phi \text{ gemm}(\_, \text{zipwith}(\ominus)) b_1 t_2) \ominus (NIL \phi s_2)) \\
= & \quad \{ \text{Definition of map} \} \\
& (\text{map}(\text{map } \text{sum})_{s_1} \phi \text{ map}(\text{map } \text{sum}) (\text{gemm}(\_, \text{zipwith}(\ominus)) b_1 t_2)) \\
& \quad \ominus (NIL \phi \text{ map}(\text{map } \text{sum})_{s_2}) \\
= & \quad \{ \text{Promotion of map, folding} \} \\
& (h_1 s_1 \phi \text{ gemm}(\_, \text{zipwith}(+))(\text{map}(\text{map } \text{sum}) b_1) (\text{map}(\text{map } \text{sum}) t_2)) \\
& \quad \ominus (NIL \phi h_1 s_2)
\end{aligned}$$

In the last formula, functions applied to  $t_1$  and  $b_1$  should be  $h_2$  and  $h_3$  respectively, which suggests us to define  $h_2$ ,  $h_3$  and  $\odot_1$  as follows.

$$\begin{aligned}
h_1 = h_2 = h_3 &= \text{map}(\text{map } \text{sum}) \\
& (s_1, t_1, b_1, r_1, l_1, tr_1, br_1, tl_1, bl_1, c_1, ro_1) \\
\odot_1 & (s_2, t_2, b_2, r_2, l_2, tr_2, br_2, tl_2, bl_2, c_2, ro_2) \\
& = (s_1 \phi \text{ gemm}(\_, \text{zipwith}(+)) b_1 t_2) \ominus (NIL \phi s_2)
\end{aligned}$$

Similarly, we can derive  $\ominus_1$  by calculating  $h_1 (x \otimes_1 y)$  as follows:

$$\begin{aligned}
& (s_1, t_1, b_1, r_1, l_1, tr_1, br_1, tl_1, bl_1, c_1, ro_1) \\
\ominus_1 & (s_2, t_2, b_2, r_2, l_2, tr_2, br_2, tl_2, bl_2, c_2, ro_2) \\
= & \text{zipwith}_4 f_s s_1 s_2 r_1 l_2 \\
& \text{where } f_s s_1 s_2 r_1 l_2 = (s_1 \phi \text{ gemm}(\_, +) r_1 l_2) \ominus (NIL \phi s_2)
\end{aligned}$$

and derive other functions and operators by doing similarly about  $\oplus_i$  and  $\otimes_i$ . Finally, we get the following.

$$h_i = \text{map}(\text{map } \text{sum}) \quad (i = 1, \dots, 9)$$

$$h_{10} = h_{11} = \text{map } \text{sum}$$

$$\text{map}(\text{map } \text{sum}) \circ \text{rects} = \pi_1 \circ (\Delta_1^{11} f'_i, \Delta_1^{11} \odot_i, \Delta_1^{11} \ominus_i)$$

**where**

$$f'_1 |a| = ||a||$$

Repeating such fusion with `map max` and `max` will yield the result shown in Figure 7. In this final program, each of the nine elements of the resulting tuple is the partial answer of its counterpart in Figure 8. Some parts are calculated with a general matrix multiplication, and others are updated with `map` and `zipwith`. For example, the element  $s_0$ , which is the solution of the maximum rectangle sum, is the maximum of the solutions of upper and lower subarray or the maximum of the solutions generated by combining partial answers of the top and the bottom rectangles as shown in Figure 4-(a). Here, the rectangles that share the same edge are combined by `zipwith(+)`. Similarly, some elements of the array  $r_0$ , which is the partial solutions of the rectangles on the right edge, are calculated with a general matrix multiplication as shown in Figure 4-(b). Here,  $(i, j)$  element in the block is the maximum of  $br_1(i, k) + br_1(k, j)$  for all  $k$ .

Provided that we divide the input array into two parts evenly, this final parallel program uses only  $O(n^3)$  addition operations as follows. For an  $n \times n$  input array, the program's cost  $T(n, n)$  satisfies the next

equation with *gemm*'s cost  $T_{gemm}(n, n)$  and some constants  $c_1$  and  $c_2$ .

$$T(n, n) = 4T(n/2, n/2) + c_1T_{gemm}(n/2, n/2) + c_2n^2$$

Since the cost of the general matrix multiplication (*gemm*) is  $O(n^3)$ , the answer of the above equation is  $T(n, n) = O(n^3)$ . This is much better than the initial one. Moreover, since the general matrix multiplication  $gemm(\uparrow, +)$  used in the program is a distance matrix multiplication, we can achieve subcubic cost with the special implementation of the  $gemm(\uparrow, +)$  used in Takaoka's algorithm <sup>(41)</sup>. However, the implementation is somewhat tricky, so that we cannot describe it with our skeletons.

#### STEP 4. OPTIMIZING INNER FUNCTIONS

For our example, we may proceed to optimize the operators and functions such as  $f_i'''$ ,  $\odot_i''$  and  $\ominus_i''$  in the program of Step 3. Since they cannot be made efficient any more, we finish our derivation of an efficient parallel program.

### 5. Implementation

In this section, we will give an efficient parallel implementation (on PC clusters) of the parallel skeletons, which are primitive operations on two-dimensional arrays defined in Section 3.1 and Section 3.2. Since a homomorphism can be specified as a composition of the reduce and map skeletons, homomorphisms have efficient parallel implementations.

Our parallel skeletons are implemented as a C++ library with MPI. We will report some experimental results, showing programs described with skeletons can be executed efficiently in parallel.

### 5.1. IMPLEMENTATION OF DATA PARALLEL SKELETONS

The four basic data parallel skeletons of `map`, `zipwith`, `reduce` and `scan` can be efficiently implemented on distributed memory systems. To illustrate this, we separate computations of a skeleton into two parts: local computations within a processor and global computations crossing processors.

For `map` skeleton, we can separate its computation as follows.

$$\begin{aligned}
 \text{map } f &= \text{map } f \circ \text{gather} \circ \text{dist } p q \\
 &= \text{map } f \circ \text{reduce}(\oplus, \phi) \circ \text{dist } p q \\
 &= \text{reduce}(\oplus, \phi) \circ \text{map}(\text{map } f) \circ \text{dist } p q \\
 &= \text{gather} \circ \text{map}(\text{map } f) \circ \text{dist } p q
 \end{aligned}$$

The last formula indicates that we can compute `map f` by distributing a two-dimensional array of the argument to the processors by `dist p q`, applying `map f` to each local array independently on each processor, and finally gathering the results onto the root processor by `gather`. Thus, for a two-dimensional array of  $n \times n$  size we can compute `map f` in  $O(n^2/P)$  parallel time, using  $P = pq$  processors and ignoring distribution and collection provided that the function  $f$  can be computed in  $O(1)$  time. This is the same also about `zipwith`.

For reduce skeleton, we can separate its computation as follows.

$$\begin{aligned}
\text{reduce}(\oplus, \otimes) &= \text{reduce}(\oplus, \otimes) \circ \text{gather} \circ \text{dist } p q \\
&= \text{reduce}(\oplus, \otimes) \circ \text{reduce}(\ominus, \phi) \circ \text{dist } p q \\
&= \text{reduce}(\oplus, \otimes) \circ \text{map}(\text{reduce}(\oplus, \otimes)) \circ \text{dist } p q
\end{aligned}$$

The last formula indicates that we can compute  $\text{reduce}(\oplus, \otimes)$  by distributing a two-dimensional array of the argument to the processors by  $\text{dist } p q$ , applying  $\text{reduce}(\oplus, \otimes)$  to each local array independently on each processor, and finally reducing the results into the root processor by  $\text{reduce}(\oplus, \otimes)$  described in the last formula. From the property of Eq. (1), the last reduction over the results of all processors can be computed by using tree-like computation in column and row directions respectively like parallel computation of reduction on one-dimensional lists. Thus, for a two-dimensional array of  $n \times n$  size we can compute  $\text{reduce}(\oplus, \otimes)$  in  $O(n^2/P + \log P)$  parallel time, using  $P = pq$  processors and ignoring distribution provided that the binary operators  $\oplus$  and  $\otimes$  can be computed in  $O(1)$  time. Note that we can also execute the tree-like computation in both of column and row directions simultaneously.

For scan skeleton, we can separate its computation as follows.

$$\begin{aligned}
\text{scan}(\oplus, \otimes) &= \text{reduce}(\oplus', \otimes') \circ \text{map } |\cdot| \circ \text{gather} \circ \text{dist } p q \\
&= \text{reduce}(\oplus', \otimes') \circ \text{map } |\cdot| \circ \text{reduce}(\ominus, \phi) \circ \text{dist } p q \\
&= \text{reduce}(\oplus', \otimes') \circ \text{map}(\text{reduce}(\oplus', \otimes') \circ \text{map } |\cdot|) \circ \text{dist } p q \\
&= \text{reduce}(\oplus', \otimes') \circ \text{map}(\text{scan}(\oplus, \otimes)) \circ \text{dist } p q \\
&= \text{gather} \circ \underline{\text{dist } p q} \circ \text{reduce}(\oplus', \otimes') \circ \text{map}(\text{scan}(\oplus, \otimes)) \circ \text{dist } p q
\end{aligned}$$

The second last formula indicates we can compute  $\text{scan}(\oplus, \otimes)$  by distributing a two-dimensional array of the argument to the processors by  $\text{dist } pq$ , applying  $\text{scan}(\oplus, \otimes)$  to each local array independently on each processor, and finally reducing the results into the root processor by  $\text{reduce}(\oplus', \otimes')$ . However, since the result of  $\text{scan}(\oplus, \otimes)$  is a two-dimensional array, we want that the last operation of computing  $\text{scan}(\oplus, \otimes)$  is  $\text{gather}$  like the case of  $\text{map } f$ . Thus, we compute underlined  $\text{dist } pq \circ \text{reduce}(\oplus', \otimes')$  instead of the last reduction  $\text{reduce}(\oplus', \otimes')$ . Although under our notation the underlined computation cannot be written in simpler form, we can compute it in sequence in column and row direction like the case of  $\text{reduce}$ . The computation in each direction can be done like those of lists <sup>(21)</sup>. From the property of Eq. (2), we can also compute  $\text{scan}(\oplus, \otimes)$  by computing  $\text{scan}_{\downarrow}(\oplus)$  after  $\text{scan}_{\rightarrow}(\otimes)$ . Note that  $\text{scan}_{\downarrow}(\oplus)$  and  $\text{scan}_{\rightarrow}(\otimes)$  can be computed in the same way of  $\text{scan}$  on lists although it performs to two or more lists simultaneously. Thus, for a two-dimensional array of  $n \times n$  size we can compute  $\text{scan}(\oplus, \otimes)$  in  $O(n^2/P + \sqrt{n^2/P} \log P)$  parallel time, using  $P = pq$  processors and ignoring distribution and collection provided that the binary operators  $\oplus$  and  $\otimes$  can be computed in  $O(1)$  time. Note that we can also execute the global computation in both of column and row directions simultaneously.

Finally, we note that gathering and re-distribution between successive calls of skeletons can be canceled when  $\text{dist } pq \circ \text{gather} = \text{id}$ . This condition is satisfied when the argument arrays of the successive calls of skeletons are distributed in the same way. That is, once argument arrays are distributed, we do not need extra communication caused by re-distribution of arrays. Since this situation occurs very often, we omit

communication cost of distribution of argument arrays in the cost of skeletons. However, we sometimes need to change distribution of argument arrays in some computations. For example, iterated matrix-vector multiplication  $x_n = Ax_{n-1}$ , which is the kernel of many important algorithms, needs such re-distribution between steps. In the computation, we distribute the matrix  $A$  row-wisely and duplicate  $x_{n-1}$  onto each processor. Each processor computes a slice of  $x_n$  and then broadcasts it to all other processors for the next iteration. In this case, the communication cost of the re-distribution (the broadcast) cannot be factored out by the condition  $\text{dist } p q \circ \text{gather} = \text{id}$ . To take the communication cost into account, we represent the communication via computation with our skeletons. To illustrate it, we formalize the example as follows.

$$\begin{aligned}
 & \text{mvmul } n \ A \ x_0 = \text{iter } n \ \text{step } x_0 \\
 & \textbf{where } \text{step} = (\text{zipwith } \text{iproduct } rA) \circ \text{duplicate} \\
 & \quad \text{duplicate} = \text{scanr}(\gg, \gg) \circ \text{scan}(\ominus, \phi) \circ \text{map } |\cdot| \\
 & \quad rA = \text{rows } A \\
 & \quad \text{rows} = \text{reduce}(\ominus, \text{zipwith}(\phi)) \circ \text{map } \|\cdot\|
 \end{aligned}$$

In each iteration step, the argument vector  $x_{n-1}$  is redistributed (broadcasted) by *duplicate*, which is a composition of skeletons, and then each copy is used to produce an inner product with a row of the matrix  $A$ , which is distributed row-wisely and named  $rA$ . This *duplicate* seems not to do actual computation but does computation of re-distribution, i.e. it copies a partial result on other processor and concatenates it to its own partial result. Since the cost of the concatenating operators (i.e.  $\ominus$  and  $\phi$ ) for arrays of  $N$  elements is considered to be  $O(N)$  in the worst

case, the cost of *duplicate* is  $O(N \log N)$  and this is equal to the cost of the usual communication. This is the idea to take the communication cost into account. Note that gathering and re-distribution between successive steps in the iteration can be canceled since the distribution of the argument vector  $x_n$  is the same at beginning of each step.

Besides *duplicate*, we can represent several communications such as broadcasting and shifting via our skeletons.

$$\begin{aligned}
 \mathit{bcast} &= \mathit{scan}(\lll, \lll) \\
 \mathit{shift}_C &= \mathit{map}(\pi_1) \circ \mathit{scan}(\ggg, \odot) \circ \mathit{map} \ \mathit{init} \\
 \mathit{shift}_R &= \mathit{map}(\pi_1) \circ \mathit{scan}(\odot, \ggg) \circ \mathit{map} \ \mathit{init} \\
 &\mathbf{where} \ \mathit{init} \ a = (\epsilon, a) \\
 &\quad (x, a) \odot (\epsilon, b) = (a, b) \\
 &\quad (x, a) \odot (y, b) = (y, b) \quad (y \neq \epsilon)
 \end{aligned}$$

The function *bcast* broadcasts the element on the root processor to the other processors. The column-shifting function *shift<sub>C</sub>* shifts each column to the right and fills the first column with special value  $\epsilon$ . The row-shifting function *shift<sub>R</sub>* shifts each row to the bottom. As illustrated above, we represent explicit communications by computations with skeletons. However, sometimes those representations have more costs than usual implementations and we need to consider special implementations. Some of such communications are discussed in the next section.

## 5.2. IMPLEMENTATION OF DATA COMMUNICATION SKELETONS

We have efficient parallel implementations for the data communication skeletons defined in Section 3.2.

Since `dist` distributes all elements of a two-dimensional array at the root processor to all other processors and `gather` does the inverse, we can compute `dist` and `gather` in  $O(n^2)$  parallel time for a two-dimensional array of  $n \times n$  size.

Although the definition of  $\text{rot}_r f$  given in Section 3 is complicated, the actual operation of  $\text{rot}_r f$  is simple. Function  $\text{rot}_r f$  merely rotates independently the  $i$ -th row by  $f i$ , and rotation of each row can be done by four parallel communications. Without losing generality we can assume that the amount of rotation  $r = f i$  satisfies  $0 < r \leq n/2$  where  $n$  is the length of the row because we just reverse the direction of rotation in the case of  $n/2 < r$ . The operations are followings: (1) making groups of  $2r$  processors from the first processor of the row (i.e.  $n/(2r)$  groups are made) and transmitting subarrays of first  $r$  processors to the rest  $r$  processors in each group, (2) considering that processors from the 0th to the  $r$ -th continue behind the last processor, making groups of  $2r$  processors from the  $r$ -th processor of the row and transmitting subarrays of first  $r$  processors to the rest  $r$  processors in each group (i.e. processors in the first  $n/(2r)$  groups have transmitted their subarrays), (3) doing the former two operations on the rest processors that have not transmitted their subarrays yet, considering the processors that have done continue behind the processors. Since more than the half processors have transmitted their subarrays by the end of the former two operations, all processors can transmit their subarrays by the end

of third operation. Thus, since the amount of one communication is  $O(n^2/P)$  for  $P$  processors,  $\text{rot}_r f$  can be executed in  $O(n^2/P)$  parallel time. Similarly,  $\text{rot}_c f$  can be executed in  $O(n^2/P)$  parallel time.

### 5.3. OPTIMIZATION BY FUSION

Although it is easy to make a parallel program by composing parallel skeletons, a simply composed skeletal program has overheads due to redundant intermediate data structures, communications and synchronizations. Thus, we perform optimization on user-written simple skeleton programs by fusing successive skeletons to eliminate redundant intermediate data structures, communications and synchronizations.

The main rules to perform optimization by fusion in our current system are as follows.

$$\begin{aligned} \text{map } f \circ \text{map } g &\Rightarrow \text{map } (f \circ g) \\ \text{reduce}(\oplus, \otimes) \circ \text{map } f &\Rightarrow (\lfloor f, \oplus, \otimes \rfloor) \\ (\lfloor f, \oplus, \otimes \rfloor) \circ \text{map } g &\Rightarrow (\lfloor f \circ g, \oplus, \otimes \rfloor) \end{aligned}$$

These rules are instances of the fusion rule of Theorem 2.3. Successive skeletons on the left hand side are fused into one skeleton on the right hand side. Then, intermediate data structures, communications and synchronizations between skeletons on the left hand side are eliminated in the right hand side. Here, a homomorphism  $(\lfloor f, \oplus, \otimes \rfloor)$  is just used as an internal representation for the optimization, and users do not need to consider homomorphisms explicitly. Since a homomorphism clearly has an efficient implementation in parallel, we use the implementation for homomorphisms produced by the optimization.

This optimization can be done automatically by the library, in the similar way to automatic optimization for lists <sup>(32)</sup>. It is our future work to formalize further rules for optimization by fusion.

#### 5.4. EXPERIMENTAL RESULTS

We implemented the parallel skeletons as part of the skeleton library *SkeTo* <sup>(31)</sup> <sup>1</sup> with C++ and MPI, and did our experiment on a cluster (distributed memory). Each of the nodes connected with Gigabit Ethernet has a CPU of Intel® Xeon®2.80GHz and 2GB memory, with Linux 2.4.21 for the OS, gcc 4.1.1 for the compiler, and mpich 1.2.7 for the MPI.

We measured execution times and speedups for the following parallel programs described by the parallel skeletons:

*Frobenius Norm*

$$fnorm = \text{reduce}(+, +) \circ \text{map } square ,$$

*Matrix Multiplication*

$$mm \text{ (composition of skeletons; see Section 3.3) ,}$$

*Maximum Rectangle Sum*

$$mrs \text{ (derived program written by skeletons; see Section 4.2) .}$$

Figures 9 through 11 show measured speedups of the programs and their optimized versions. Optimized programs are named XXX\_opt. The speedup is a ratio of running time of the program on  $p$  processors to that of hand-written sequential program on one processor, without first distribution of arrays. Tables I through III show measured running

<sup>1</sup> It is available at <http://www.ipl.t.u-tokyo.ac.jp/sketo/>

times and speedups. Besides speedups with respect to hand-written sequential programs, speedups with respect to the skeleton programs on one processor are listed in the tables. The inputs are an  $8000 \times 8000$  matrix for *fnorm*, a  $3000 \times 3000$  matrix for *mm*, and a  $1000 \times 1000$  matrix for *mrs*. The running time of *mrs* is relatively big because its operators in its calculation are more complicated than those of the others.

The result shows programs described with skeletons can be executed efficiently in parallel, and proves the success of our framework. Both simply composed skeleton programs and optimized programs achieve good speedups. The optimized programs achieve better running time than the simple programs. Since *mrs* requires a large amount of communication data and nested parallelism, speedup of *mrs* on 64 processors for a  $1000 \times 1000$  matrix is not so good.

To examine sequential performances of our skeleton programs, we implemented sequential programs for the above programs. We compared their calculation times with those of the skeleton programs on one processor. The program *fnorm* simply written with our skeletons is five times slower than the hand-made sequential program. This is caused by overhead due to creation of intermediate data between `map square` and `reduce(+, +)`. Thus, the program optimized by fusion achieves the same running time as the sequential program due to elimination of the intermediate data. The program *mm* written with our skeletons is two times slower than the hand-made sequential program. This is mainly caused by overhead of methods for accessing elements of matrices, and some compilers can eliminate this overhead. Thus, the program optimized by fusion achieves better running time than the simple program, but it is

still 1.6 times slower than the sequential program. Since operators in the calculation of *mrs* are complicated, the overhead of skeletons are relatively small. Thus, the running times of sequential program, simple program and optimized program are almost the same.

Finally, we list part of the C++ code of *mm* written with the skeleton library in Figure 12, to give a concrete impression of the conciseness our library provides.

## 6. Related Works

Besides the related work as in the introduction, our work is closely related with the active researches on matrix representation for parallel computations and the compositional approach to parallel program development.

### RECURSIVE MATRIX REPRESENTATIONS

Wise et al. <sup>(42)</sup> proposed representation of a two-dimensional array by a quadtree, i.e. a two-dimensional array recursively constructed by four small sub-arrays of the same size. This representation is suitable for describing recursive blocked algorithms <sup>(15)</sup>, which can provide better performance than existing algorithms for some matrix computations such as LU and QR factorizations <sup>(18,43)</sup>. However, the quadtree representation requires the size of two-dimensional arrays to be the power of two. Moreover, once a two-dimensional array is represented by a quadtree, we cannot reblock the array by restructuring the quadtree, which would

prevent us from developing more parallelism in the recursive blocked algorithms on them.

Bikshandi et al. <sup>(5)</sup> proposed representation of a two-dimensional array by a hierarchically tiled array (HTA). An HTA is an array partitioned into tiles, and these tiles can be either conventional arrays or lower level HTAs. The outermost tiles are distributed across processors for parallelism and the inner tiles are utilized for locality. In the HTA programming, users are allowed to use recursive index accessing according to the structure of HTAs, so that they can easily transform conventional programs onto HTA programs. Communication and synchronization are explicitly expressed by index accessing to remote tiles. Thus HTA programs can control relatively low-level parallelism and can be efficient implementation. However, it is not presented how to derive efficient HTA programs. We think it is good that we derive an efficient algorithm on the abide-tree then implement it on HTAs.

A more natural representation of a two-dimensional array is to use nested one-dimensional arrays (lists) <sup>(7,29,38)</sup>. The advantage is that many results developed for lists can be reused. However, this representation imposes much restriction on the access order of elements.

The abide-tree representation, as used in this paper, was first proposed by Bird <sup>(7)</sup>, as an extension of one-dimensional join lists. However, the focus there is on derivation of sequential programs for manipulating two-dimensional arrays, and there is little study on the framework for developing efficient parallel programs. Our work provides a good complement.

This work was greatly inspired by the success of compositional (skeletal) parallel programming on one-dimensional arrays (lists) <sup>(35)</sup>, and our initial motivation was to import the results so far to two-dimensional arrays. This turns out to be more difficult than we had expected.

Compositional parallel Programming using Bird-Meertens Formalism (BMF) has been attracting many researchers. The initial BMF <sup>(6)</sup> was designed as a calculus for deriving (sequential) efficient programs on lists. Skillicorn <sup>(37)</sup> showed that BMF could also provide an architecture independent parallel model for parallel programming because a small fixed set of higher order functions (skeletons) in BMF such as `map` and `reduce` can be mapped efficiently to a wide range of parallel architectures.

Systematic programming methods have actively been studied in the framework of skeletal (compositional) parallel programming on lists. The diffusion theorem <sup>(27)</sup> gives a powerful method to obtain suitable composition of skeletons for a program recursively defined on lists and trees. Chin et al. <sup>(10,26)</sup> have studied a systematic method to derive an associative operator that plays an important role in parallelization, based on which Xu et al. <sup>(44)</sup> build an automatic derivation system for parallelizing recursive linear functions with normalization rules.

Our parallel skeletons can be seen as an instance of design-patterns in object-oriented programmings. The difference between our approach and usual design-pattern approaches is that we use a small number of skeletons carefully designed rather than heedlessly increasing the number of patterns. This enables us to make a clear automatic optimization

mechanism. We also give a methodology to derive an efficient program described with skeletons.

APL <sup>(4,17,28)</sup> is a pioneer language that supports operators to manipulate arrays, namely array operators. APL's array operators are extensions of scalar operations and functions to multi-dimensional arrays in an element-wise manner, manipulations of the layout of arrays such as shift and rotate, and reduction operators to collapse arrays with binary operators. More complex computations on arrays are constructed by composition of these array operators. The ideas of APL and our skeletal approach are the same in the sense that both provide users with a set of basic patterns of array computations and let users make sophisticated computations by compositions of these basic patterns. Also, both array operators and skeletons conceal complicated parallelism from users. Main difference between APL and our skeletal approach is as follows. We give a systematic methodology to derive (make) correct, safe and efficient programs, i.e. how to derive efficient programs from correct and safe programs that are not so efficient. Our skeletons treat 2D arrays as '2D' in reductions since they can use two binary operators for horizontal direction and vertical direction, while treatment of multidimensional arrays in reductions of APL is rather like '1D' because they use only one operator in one reduction. Our skeletons are designed to consider derivations and optimizations for parallel programs of compositions of skeletons based on the mathematical theory.

## 7. Conclusion

In this paper, we propose a compositional framework that allows users, even with little knowledge about parallel machines, to describe safe and efficient parallel computation over two-dimensional arrays easily. In our framework, two-dimensional arrays are represented by the abide-tree that supports systematic development of parallel programs and architecture-independent implementation, and programmers can easily build up a complicated parallel system by defining basic components recursively, putting components compositionally, and improving efficiency systematically. The power of our approach is seen from the nontrivial programming examples of matrix multiplication and QR decomposition, and a successful derivation of an involved efficient parallel program for the maximum rectangle sum problem <sup>(24)</sup>. A demonstration of an efficient implementation of basic computation skeletons (in C++ and MPI) on distributed PC clusters guarantees that programs composed by these parallel skeletons can be efficiently executed in parallel.

Currently, we have not succeeded in developing efficient and correct parallel programs with parallel skeletons from their recursive specifications such as nested recursive functions on nested lists and divide-and-conquer algorithm on quad-trees. Thus, it is our future work to construct more powerful theories for a systematic programming methodology that supports such derivations, and to introduce control skeletons such as divide-and-conquer skeleton if needed. It is also our future work to study an automatic optimization mechanism. The mechanism should be able to eliminate inefficiency due to compositional or nested

uses of parallel skeletons in parallel programs. Moreover, it should be able to introduce low-level parallelism that currently our skeletons cannot describe. Some of the optimizations will be performed based on the properties of operators in the skeletons. Finally, we mention about higher-dimensional arrays. To extend our framework to higher-dimensional arrays, we introduce constructors for new dimensions and the condition that any pair of the constructors satisfies the abide-property. Then, definitions of parallel skeletons and programming methodology for higher-dimensional arrays are almost the same as two-dimensional.

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$$\begin{array}{l}
\text{map } f \begin{pmatrix} x_{11} & x_{12} & \cdots & x_{1n} \\ x_{21} & x_{22} & \cdots & x_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ x_{m1} & x_{m2} & \cdots & x_{mn} \end{pmatrix} = \begin{pmatrix} f \ x_{11} & f \ x_{12} & \cdots & f \ x_{1n} \\ f \ x_{21} & f \ x_{22} & \cdots & f \ x_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ f \ x_{m1} & f \ x_{m2} & \cdots & f \ x_{mn} \end{pmatrix} \\
\\
\text{reduce}(\oplus, \otimes) \begin{pmatrix} x_{11} & x_{12} & \cdots & x_{1n} \\ x_{21} & x_{22} & \cdots & x_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ x_{m1} & x_{m2} & \cdots & x_{mn} \end{pmatrix} = \begin{pmatrix} (x_{11} \otimes x_{12} \otimes \cdots \otimes x_{1n}) \oplus \\ (x_{21} \otimes x_{22} \otimes \cdots \otimes x_{2n}) \oplus \\ \vdots \\ (x_{m1} \otimes x_{m2} \otimes \cdots \otimes x_{mn}) \end{pmatrix} \\
\\
\text{zipwith } f \begin{pmatrix} x_{11} & x_{12} & \cdots & x_{1n} \\ x_{21} & x_{22} & \cdots & x_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ x_{m1} & x_{m2} & \cdots & x_{mn} \end{pmatrix} \begin{pmatrix} y_{11} & y_{12} & \cdots & y_{1n} \\ y_{21} & y_{22} & \cdots & y_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ y_{m1} & y_{m2} & \cdots & y_{mn} \end{pmatrix} \\
= \begin{pmatrix} f \ x_{11} \ y_{11} & f \ x_{12} \ y_{12} & \cdots & f \ x_{1n} \ y_{1n} \\ f \ x_{21} \ y_{21} & f \ x_{22} \ y_{22} & \cdots & f \ x_{2n} \ y_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ f \ x_{m1} \ y_{m1} & f \ x_{m2} \ y_{m2} & \cdots & f \ x_{mn} \ y_{mn} \end{pmatrix} \\
\\
\text{scan}(\oplus, \otimes) \begin{pmatrix} x_{11} & x_{12} & \cdots & x_{1n} \\ x_{21} & x_{22} & \cdots & x_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ x_{m1} & x_{m2} & \cdots & x_{mn} \end{pmatrix} = \begin{pmatrix} y_{11} & y_{12} & \cdots & y_{1n} \\ y_{21} & y_{22} & \cdots & y_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ y_{m1} & y_{m2} & \cdots & y_{mn} \end{pmatrix} \\
\text{where } y_{ij} = \begin{pmatrix} (x_{11} \otimes x_{12} \otimes \cdots \otimes x_{1j}) \oplus \\ (x_{21} \otimes x_{22} \otimes \cdots \otimes x_{2j}) \oplus \\ \vdots \\ (x_{i1} \otimes x_{i2} \otimes \cdots \otimes x_{ij}) \end{pmatrix}
\end{array}$$

Figure 1. Intuitive Definition of Four Primitive Skeletons on Two-Dimensional Arrays

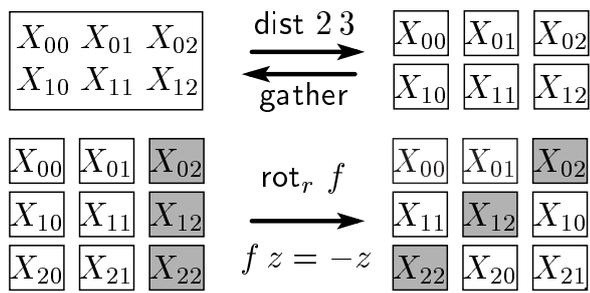
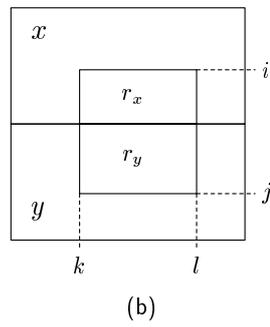


Figure 2. An Image of Communication Skeletons (each rectangle corresponds to each processor;  $X_{ij}$  represents a subarray.)

$$\text{rects} \left( \begin{array}{c} \overbrace{\left( \begin{array}{c} m_x \\ \boxed{x} \end{array} \right)}^n \\ \overbrace{\left( \begin{array}{c} m_x \\ \boxed{y} \end{array} \right)} \end{array} \right) = \begin{array}{c} \xrightarrow{j} \\ \left( \begin{array}{cc} \overbrace{\left( \begin{array}{c} m_x \\ \text{rects } x \end{array} \right)} & \boxed{Z} \\ & \overbrace{\left( \begin{array}{c} m_y \\ \text{rects } y \end{array} \right)} \\ \end{array} \right) \downarrow i
 \end{array}$$

(a)



$$\boxed{Z} = \begin{array}{c} \boxed{\phantom{Z}} \\ \text{bottoms } x \end{array} \times \boxed{\text{tops } y}$$

(c)

Figure 3. Recursive Computation of *rects*

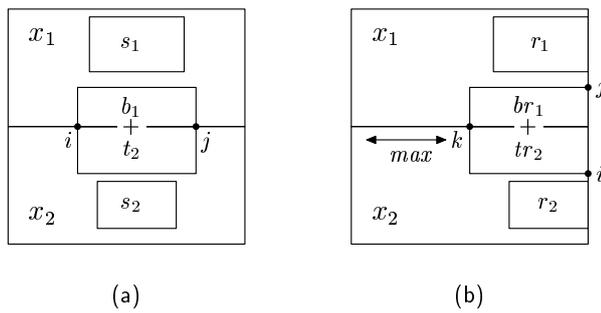


Figure 4. Computation of the Operator for Derived  $mrs$

---

```

tops |a|      = |||a|||
tops (x ⊗ y) = tops x ⊕ map (zipwith(⊗) (cols' x)) (tops y)
tops (x ⊕ y) = zipwith4 ft (tops x) (tops y) (toprights x) (toplefts y)
               where ft t1 t2 tr1 tl2 = (t1 ⊕ gemm (⊖, ⊕) tr1 tl2) ⊗ (NIL ⊕ t2)

bottoms |a|   = |||a|||
bottoms (x ⊗ y) = map (λz → zipwith(⊗) z (cols' y)) (bottoms x) ⊗ bottoms y
bottoms (x ⊕ y) = zipwith4 fb (bottoms x) (bottoms y) (bottomrights x) (bottomlefts y)
               where fb b1 b2 br1 bl2 = (b1 ⊕ gemm (⊖, ⊕) br1 bl2) ⊗ (NIL ⊕ b2)

rightss |a|   = |||a|||
rightss (x ⊗ y) = (rightss x ⊕ gemm (⊖, zipwith(⊗)) (bottomrights x) (toprights y))
                  ⊗ (NIL ⊕ rightss y)
rightss (x ⊕ y) = zipwith3 fr (rightss x) (rightss y) (rows' y)
               where fr r1 r2 ro2 = map (⊕ ro2) r1 ⊗ r2

leftss |a|    = |||a|||
leftss (x ⊗ y) = (leftss x ⊕ gemm (⊖, zipwith(⊗)) (bottomlefts x) (toplefts y)) ⊗ (NIL ⊕ leftss y)
leftss (x ⊕ y) = zipwith3 fl (leftss x) (leftss y) (rows' x)
               where fl l1 l2 ro1 = l1 ⊕ map (ro1 ⊕) l2

toprights |a| = |||a|||
toprights (x ⊗ y) = toprights x ⊕ map (zipwith(⊗) (right' (toprights x))) (toprights y)
toprights (x ⊕ y) = zipwith ftr (toprights x) (toprights y)
               where ftr tr1 tr2 = map (⊕ top' tr2 ⊕) tr1 ⊗ tr2

bottomrights |a| = |||a|||
bottomrights (x ⊗ y) = map (λz → zipwith(⊗) z (top' (bottomrights y))) (bottomrights x)
                  ⊗ bottomrights y
bottomrights (x ⊕ y) = zipwith fbr (bottomrights x) (bottomrights y)
               where fbr br1 br2 = map (⊕ top' br2 ⊕) br1 ⊗ br2

toplefts |a|   = |||a|||
toplefts (x ⊗ y) = toplefts x ⊕ map (zipwith(⊗) (right' (toplefts x))) (toplefts y)
toplefts (x ⊕ y) = zipwith ftl (toplefts x) (toplefts y)
               where ftl tl1 tl2 = tl1 ⊕ map (right' tl1 ⊕) tl2

bottomlefts |a| = |||a|||
bottomlefts (x ⊗ y) = map (λz → zipwith(⊗) z (top' (bottomlefts y))) (bottomlefts x)
                  ⊗ bottomlefts y
bottomlefts (x ⊕ y) = zipwith fbl (bottomlefts x) (bottomlefts y)
               where fbl bl1 bl2 = bl1 ⊕ map (right' bl1 ⊕) bl2

cols' |a|     = ||a||
cols' (x ⊗ y) = zipwith(⊗) (cols' x) (cols' y)
cols' (x ⊕ y) = (cols' x ⊕ gemm (⊖, ⊕) (right (cols' x)) (top (cols' y))) ⊗ (NIL ⊕ cols' y)

rows' |a|     = ||a||
rows' (x ⊗ y) = (rows' x ⊕ gemm (⊖, ⊗) (right (rows' x)) (top (rows' y))) ⊗ (NIL ⊕ rows' y)
rows' (x ⊕ y) = zipwith(⊕) (rows' x) (rows' y)

top          = reduce(⊕, ⊕) ∘ map |·|
bottom      = reduce(⊗, ⊕) ∘ map |·|
right       = reduce(⊗, ⊗) ∘ map |·|
left        = reduce(⊗, ⊕) ∘ map |·|
top'        = the ∘ top
bottom'     = the ∘ bottom
right'      = the ∘ right
left'       = the ∘ left

```

---

Figure 5. Extra Functions for *rects*

$$\begin{array}{l}
\mathit{rects} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} (a) & (a \ b) \\ (c) & (c \ d) \end{pmatrix} \begin{pmatrix} \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} & \begin{pmatrix} (c) & (c \ d) \end{pmatrix} \end{pmatrix} \\
\mathit{tops} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} (a) & (a \ b) \\ (c) & (c \ d) \end{pmatrix} \begin{pmatrix} \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} & \begin{pmatrix} (c) & (c \ d) \end{pmatrix} \end{pmatrix} \quad \mathit{bottoms} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} & \begin{pmatrix} (c) & (c \ d) \end{pmatrix} \end{pmatrix} \\
\mathit{rights} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} (a \ b) \\ (b) \end{pmatrix} \begin{pmatrix} \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} \\ \begin{pmatrix} (c \ d) \\ (d) \end{pmatrix} \end{pmatrix} \quad \mathit{lefts} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} (a) & (a \ b) \\ \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} (c) & (c \ d) \end{pmatrix} \end{pmatrix} \\
\mathit{toprights} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} (a \ b) \\ (b) \end{pmatrix} \begin{pmatrix} \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} \\ \begin{pmatrix} (c \ d) \\ (d) \end{pmatrix} \end{pmatrix} \quad \mathit{bottomrights} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} \\ \begin{pmatrix} (c \ d) \\ (d) \end{pmatrix} \end{pmatrix} \\
\mathit{toplefts} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} (a) & (a \ b) \\ \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} (c) & (c \ d) \end{pmatrix} \end{pmatrix} \quad \mathit{bottomlefts} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} (c) & (c \ d) \end{pmatrix} \end{pmatrix} \\
\mathit{cols} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} a \\ c \end{pmatrix} & \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} \end{pmatrix} \quad \mathit{rows} \begin{pmatrix} a & b \\ c & d \end{pmatrix} = \left( \begin{pmatrix} a \ b \\ c \ d \end{pmatrix} \\ \begin{pmatrix} b \\ d \end{pmatrix} \end{pmatrix}
\end{array}$$

Figure 6. Examples of Extra Functions for *rects*

---


$$mrs = \pi_1 \circ (\Delta_1^{11} f_i''', \Delta_1^{11} \odot_i'', \Delta_1^{11} \ominus_i'')$$

**where**

$$\begin{aligned} (\Delta_1^{11} f_i''') |a| &= (a, |a|, |a|, |a|, |a|, |a|, |a|, |a|, |a|, |a|, |a|) \\ (s_1, t_1, b_1, r_1, l_1, tr_1, br_1, tl_1, bl_1, c_1, ro_1) \\ (\Delta_1^{11} \odot_i'') (s_2, t_2, b_2, r_2, l_2, tr_2, br_2, tl_2, bl_2, c_2, ro_2) \\ &= (s_0, t_0, b_0, r_0, l_0, tr_0, br_0, tl_0, bl_0, c_0, ro_0) \end{aligned}$$

**where**

$$\begin{aligned} s_0 &= (s_1 \uparrow \max(\text{zipwith}(+) b_1 t_2) \uparrow s_2) \\ t_0 &= \text{zipwith}_3 f_t t_1 c_1 t_2 \\ &\quad \text{where } f_t t_1 c_1 t_2 = t_1 \uparrow (c_1 + t_2) \\ b_0 &= \text{zipwith}_3 f_b b_1 c_2 b_2 \\ &\quad \text{where } f_b b_1 c_2 b_2 = (b_1 + c_2) \uparrow b_2 \\ r_0 &= (r_1 \phi \text{ gemm } (\uparrow, +) (tr br_1) tr_2) \ominus (NIL \phi r_2) \\ l_0 &= (l_1 \phi \text{ gemm } (\uparrow, +) bl_1 (tr tl_2)) \ominus (NIL \phi l_2) \\ tr_0 &= tr_1 \phi \text{ map}_c (\text{zipwith}(+) (right tr_1)) tr_2 \\ br_0 &= \text{map}_c (\text{zipwith}(+) (left br_2)) br_1 \phi br_2 \\ tl_0 &= tl_1 \ominus \text{map}_r (\text{zipwith}(+) (bottom tl_1)) tl_2 \\ bl_0 &= \text{map}_r (\text{zipwith}(+) (top bl_2)) bl_1 \ominus bl_2 \\ c_0 &= \text{zipwith}(+) c_1 c_2 \\ ro_0 &= (ro_1 \phi \text{ gemm } (\_, +) (right ro_1) (top ro_2)) \ominus (NIL \phi ro_2) \end{aligned}$$

$$\begin{aligned} (s_1, t_1, b_1, r_1, l_1, tr_1, br_1, tl_1, bl_1, c_1, ro_1) \\ (\Delta_1^{11} \ominus_i'') (s_2, t_2, b_2, r_2, l_2, tr_2, br_2, tl_2, bl_2, c_2, ro_2) \\ &= (s_0, t_0, b_0, r_0, l_0, tr_0, br_0, tl_0, bl_0, c_0, ro_0) \end{aligned}$$

**where**

$$\begin{aligned} s_0 &= s_1 \uparrow \max(\text{zipwith}(+) r_1 l_2) \uparrow s_2 \\ t_0 &= (t_1 \phi \text{ gemm } (\uparrow, +) tr_1 tl_2) \ominus (NIL \phi t_2) \\ b_0 &= (b_1 \phi \text{ gemm } (\uparrow, +) br_1 bl_2) \ominus (NIL \phi b_2) \\ r_0 &= \text{zipwith}_3 f_r r_1 r_2 ro_2 \\ &\quad \text{where } f_r r_1 r_2 ro_2 = (r_1 + ro_2) \uparrow r_2 \\ l_0 &= \text{zipwith}_3 f_l l_1 l_2 ro_1 \\ &\quad \text{where } f_l l_1 l_2 ro_1 = l_1 \uparrow (ro_1 + l_2) \\ tr_0 &= \text{map}_r (\text{zipwith}(+) (top tr_2)) tr_1 \ominus tr_2 \\ br_0 &= \text{map}_r (\text{zipwith}(+) (top br_2)) br_1 \ominus br_2 \\ tl_0 &= tl_1 \phi \text{ map}_c (\text{zipwith}(+) (right tl_1)) tl_2 \\ bl_0 &= bl_1 \phi \text{ map}_c (\text{zipwith}(+) (right bl_1)) bl_2 \\ c_0 &= (c_1 \phi \text{ gemm } (\_, +) (right c_1) (top c_2)) \ominus (NIL \phi c_2) \\ ro_0 &= \text{zipwith}(+) ro_1 ro_2 \end{aligned}$$


---

Figure 7. Derived Efficient Program of Maximum Rectangle Sum

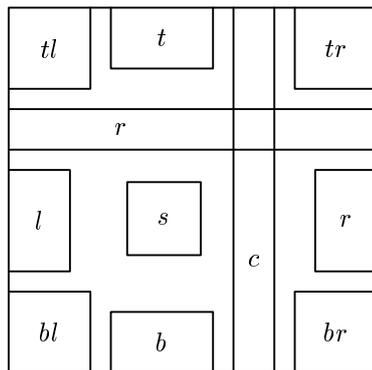


Figure 8. Corresponding Parts of Elements in the Tuple

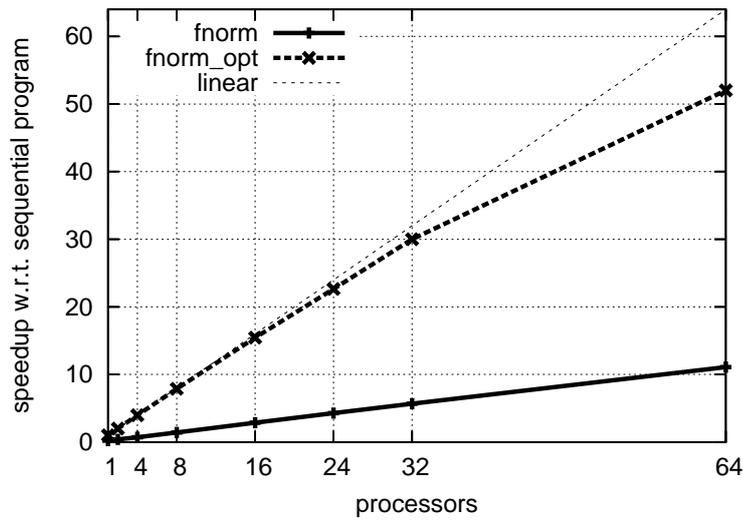


Figure 9. Speedup of F-norm

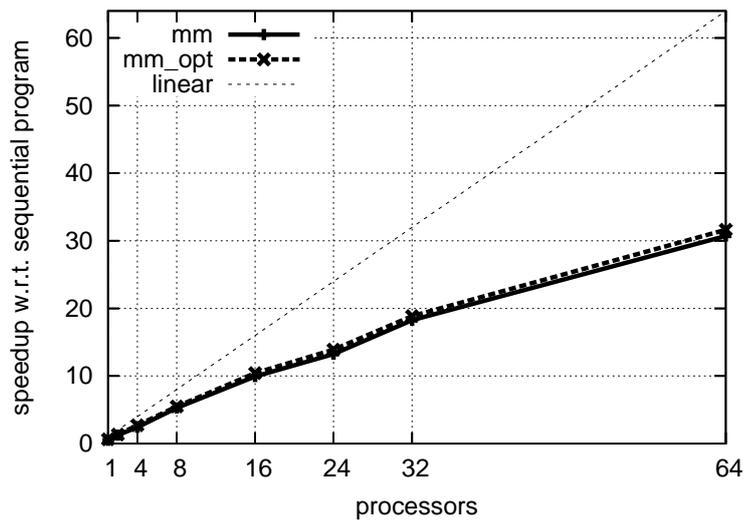


Figure 10. Speedup of Matrix Multiplication

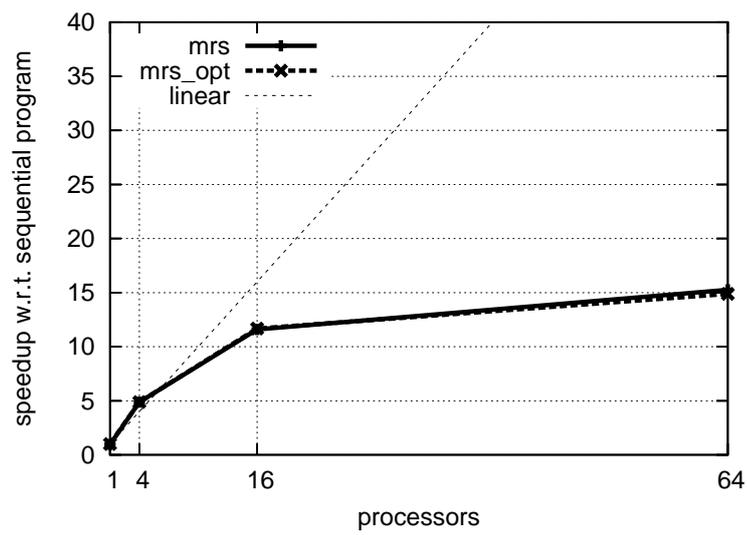


Figure 11. Speedup of Maximum Rectangle Sum

---

```
template <class C, class A, class B>
void mm(dist_matrix<C> &Z2, const dist_matrix<A> &X2, const dist_matrix<B> &Y2)
{
    dist_matrix < matrix < int > > *A2;
    dist_matrix < matrix < int > > *B2;
    A2 = all_rows2(X2);
    B2 = all_cols2(Y2);
    m_skeletons::map_i(Tri< matrix <B> >(), *B2);
    m_skeletons::zipwith(Iprod<C>(), *A2, *B2, Z2);
    delete B2;
    delete A2;
}
```

---

*Figure 12.* C++ Code of Matrix Multiplication by Parallel Skeletons

Table I. Experimental Results of F-norm for  $8000 \times 8000$  Matrices

#processors	1	2	4	8	16	24	32	64
<b>fnorm</b>								
time (s)	1.40	0.70	0.39	0.19	0.097	0.065	0.049	0.025
speedup	1.00	1.99	3.59	7.21	14.45	21.55	28.65	55.86
speedup <sup>s</sup>	0.20	0.40	0.71	1.43	2.87	4.27	5.68	11.08
<b>fnorm_opt</b>								
time (s)	0.28	0.14	0.070	0.035	0.018	0.012	0.009	0.005
speedup	1.00	1.99	3.98	7.88	15.46	22.65	30.00	52.01
speedup <sup>s</sup>	1.00	1.99	3.98	7.88	15.46	22.65	30.00	52.01

<sup>†</sup> speedup<sup>s</sup> is a speedup with respect to hand-written sequential program.

Table II. Experimental Results of Matrix Multiplication for  $3000 \times 3000$  Matrices

#processors	1	2	4	8	16	24	32	64
<b>mm</b>								
time (s)	247.02	96.26	49.26	22.66	12.09	9.05	6.56	3.91
speedup	1.00	2.57	5.02	10.90	20.44	27.30	37.64	63.23
speedup <sup>s</sup>	0.49	1.25	2.43	5.29	9.92	13.25	18.27	30.70
<b>mm_opt</b>								
time (s)	193.34	89.86	44.17	21.89	11.51	8.64	6.36	3.79
speedup	1.00	2.15	4.38	8.83	16.80	22.39	30.42	50.99
speedup <sup>s</sup>	0.62	1.33	2.71	5.48	10.42	13.89	18.87	31.63

<sup>†</sup> speedup<sup>s</sup> is a speedup with respect to hand-written sequential program.

Table III. Experimental Results of Maximum Rectangle Sum for  $1000 \times 1000$  Matrices

#processors	1	4	16	64
<b>mrs</b>				
time (s)	157.01	31.04	13.05	9.92
speedup	1.00	5.06	12.03	15.84
speedup <sup>s</sup>	0.96	4.88	11.59	15.26
<b>mrs_opt</b>				
time (s)	151.32	31.09	12.97	10.17
speedup	1.00	4.87	11.67	14.88
speedup <sup>s</sup>	1.00	4.87	11.67	14.88

<sup>†</sup> speedup<sup>s</sup> is a speedup with respect to hand-written sequential program.