Simplifying Multiple-Statement Reductions with the Polyhedral Model

CAMBRIDGE YANG, MIT CSAIL
ERIC ATKINSON, MIT CSAIL
MICHAEL CARBIN, MIT CSAIL

A Reduction – an accumulation over a set of values, using an associative and commutative operator – is a common computation in many numerical computations, including scientific computations, machine learning, computer vision, and financial analytics.

Contemporary polyhedral-based compilation techniques make it possible to optimize reductions, such as prefix sum, in which each component of the reduction’s output potentially shares computation with another component in the reduction. Therefore an optimizing compiler can identify the computation shared between multiple components and generate code that computes the shared computation only once.

These techniques, however, do not support reductions that – when phrased in the language of the polyhedral model – span multiple statements. In such cases, existing approaches can generate incorrect code that violates the data dependencies of the original, unoptimized program.

In this work, we identify and formalize the multiple/statement reduction problem as a bilinear optimization problem. We present a heuristic optimization algorithm for these reductions, and we demonstrate that the algorithm provides optimal complexity for a set of benchmark programs from the literature on probabilistic inference algorithms, whose performance critically relies on simplifying these reductions. Specifically, the complexities for 10 of the 11 programs improve significantly by factors at least of the sizes of the input data, which are in the range of $10^4$ to $10^6$ for typical real application inputs. We also confirm the significance of the improvement by showing that the speedups in wall-clock time range from 1.1x to over $10^7$x.

1 INTRODUCTION

A reduction – an accumulation over a set of values, using an associative and commutative operator – is a common computation in many numerical computations, including scientific computations, machine learning, computer vision, and financial analytics.

For example, consider the prefix sum (PS) defined mathematically by Equation (1): the value at each index $i$ of the array $B$ is the summation of values at indices $j$ before and up to $i$ of array $A$.

Listing 1 presents a direct translation of Equation (1) to an imperative language with loops. The complexity of Listing 1 is $O(N^2)$: $O(N)$ for iterating over "$i$" and $O(N)$ for the summation over $j$.

\[
B[i] = \sum_{j=0}^{j \leq i} A[j] \quad \forall i, 0 \leq i < N \tag{1}
\]

```
1 // B is array of ints initialized to all 0
2 for(i = 0; i < N; i++)
3    for(j = 0; j<=i; j++)
4        B[i] += A[j]
```

Listing 1. Naive PS

```
1 // B is array of ints initialized to all 0
2    B[0] = A[0]
3 for(i = 1; i < N; i++)
4    B[i] = B[i-1] + A[i]
```

Listing 2. Optimized PS

Optimized Reductions. Listing 2 presents a more efficient computation implementation of PS. The complexity of the implementation in Listing 2 is $O(N)$, which is a linear speedup over the naive implementation Listing 1. The implementation achieves this speedup by exploiting the fact that consecutive iterations of the loop overlap in their computations. Specifically, for any pair of consecutive iterations, the latter iteration includes the entirety of the former iteration’s computation. Therefore, that shared computation only needs to be computed once.
Gautam and Rajopadhye [2006] in the polyhedral model community formalized the above optimizing transformation under an array equational language that supports reductions as a first class operation [Yuki et al. 2013]. They also proposed a set of techniques called Simplifying Reductions (SR). The core of SR is called Simplification Transformation (ST). At a high level, ST is a transformation in array equational language. ST takes in a pointer to a statement that is a reduction (e.g. Equation (1)) and a directed vector along which the reduction’s body (e.g. $A[j]$) presents reuse as inputs: here reuse means that the reduction’s body evaluates to the same value along the direction of the vector. Given the inputs, ST transforms the statement in consideration into a set of statements that together is semantically equivalent to the original statement, but exploits the reuse vector to reduce complexity. For example, given Equation (1) and a reuse vector $[1, 0]^T$, which satisfies that changing $i$ to $i + 1$ and $j$ to $j + 0$ (i.e. not changing $j$) does not change the evaluation of $A[j]$, ST outputs Equation (2). Translating from Equation (2) to an imperative language with loops then produces Listing 2.

$$B[0] = A[0]$$
$$B[i] = B[i - 1] + A[i] \quad \forall i, 1 \leq i < N \quad (2a)$$

Note that for one application of ST there are usually infinitely many choices for directions that present reuse. For instance, any vector $[c, 0]^T$ with constant $c$ is a valid choice for the reuse vector for Equation (1), since they all satisfy that changing from $i$ to $i + c$ and not changing $j$ does not change the evaluation of $A[j]$. As a concrete example, applying ST to Equation (1) with direction $[-1, 0]^T$ produces Equation (3). Instead of initializing $B[0]$ and computing $B[i]$s from lower indices to higher indices (i.e. left to right) as in Equation (2), Equation (3) initializes $B[N - 1]$ and computes $B[i]$s from higher to lower indices (i.e. right to left). Complexity of the translation of Equation (3) to an imperative language with loops (Listing 7 in Appendix A) is also $O(N)$.

$$B[N - 1] = \sum_{j=0}^{j<N} A[j]$$
$$B[i] = B[i + 1] - A[i] \quad \forall i, 0 \leq i < N - 1 \quad (3a)$$

Multiple Statement Reductions. However, the SR framework, including ST, proposed by Gautam and Rajopadhye [2006] is restricted to optimizing one single reduction at a time, and it does not consider multiple inter-dependent statements. This is problematic because 1) ST application introduces new dependencies, and 2) the new dependencies introduced by STs together with existing dependencies of the input program may form dependency cycle(s) in the resultant program. To see 1), Equation (2b) introduces the dependency from $B[i]$ to $B[i - 1]$, i.e., $B[i]$ must be computed after the $B[i - 1]$ for any $i \in [1, N)$. To see 2), consider Equation (4a): here we extended Equation (4a) (same as Equation (1)) by Equation (4b) and obtained a program with multiple statements. If we apply ST to Equation (4a) with the reuse vector $[-1, 0]^T$, we will get the program consisting of three statements: Equations (3a), (3b) and (4b), which contains dependency cycles. For example, using “$\stackrel{eq.}{\longrightarrow}$” to mean a dependency induced by statement eq, we note that the path $B[N - 1] \stackrel{eq. (3a)}{\longrightarrow} A[N - 1] \stackrel{eq. (4b)}{\longrightarrow} B[N - 2] \stackrel{eq. (3b)}{\longrightarrow} B[N - 1]$ forms a cycle.

$$B[i] = \sum_{j=0}^{j\leq i} A[j] \quad \forall i, 0 \leq i < N \quad (4a)$$
$$A[i + 1] = f(B[i]) \quad \forall i, 0 \leq i < N - 1 \quad (4b)$$
On the other hand, if we apply ST to Equation (4a) with reuse vector $[1, 0]^T$, we will get the program consisting of three statements: Equations (2a), (2b) and (4b), which is a valid program without any dependency cycle. Listing 3 presents a translation of this program to an imperative language with loops, and it correctly computes array $A$ and $B$ with complexity $O(N)$.

1. $B[0] = A[0]$
2. for $(i = 1; i < N; i++)$
3. $B[i] = B[i-1] + A[i]$
4. $A[i+1] = f(B[i])$

Listing 3. Optimized PS with multiple statements

In summary to the above observations, the key challenge of optimizing multiple inter-dependent statements with reductions is to consolidate ST with dependency satisfaction.

**Approach.** In this work, we term the pattern in Equation (4) a *multiple-statement* reduction. We present a new technique to automatically optimize multiple-statement reductions while soundly handling inter-statements dependencies and therefore can automatically generate the code in Listing 3. The key idea behind our approach is that our heuristic algorithm uses the original program’s affine schedule as a guide to choose among the multiple choices that can be made during the optimization process. One of our key results is that we show that even though the algorithm does not consider other viable choices during optimization, given an affine schedule of the original program and all left-hand-side arrays of reductions, the algorithm is still optimal for reductions with operators that have inverses.

**Applications.** Simplifying Reductions is a classic problem in the compiler optimization literature and it has reemerged as a primary concern for modern applications. In this work, we study a suite of 11 probabilistic inference algorithms that, with the rise of data science, artificial intelligence, and machine learning, have been established as widely studied and used algorithms across computer vision, physics, and medicine. We demonstrate that multiple-statement reductions exist in these algorithms’ natural, mathematical specifications. Moreover, delivering efficient implementations of these algorithms by hand – as is current practice – requires solving the multiple-statement reduction problem by hand, which is a tedious and error-prone endeavor. Our approach shows that it is possible to automatically generate optimized efficient algorithms from the mathematical specifications alone.

**Contributions.** In this work, we present the following contributions:

- We identify the problem of multiple-statement reductions which was not addressed in Simplifying Reductions [Gautam and Rajopadhye 2006] where only a single statement was considered. We illustrate the importance of this problem through motivating real application examples.
- We formalize the task of optimizing a multiple-statement reduction by combining the insights of the Simplifying Reduction framework with insights from ILP scheduling [Pouchet et al. 2011]. We formulate a specification of the problem as a integer bilinear program.
- We propose a heuristic algorithm to solve the above optimization problem.
- We evaluate our proposed method on benchmark suites consisting of standard probabilistic inference algorithms and probabilistic models. Our results show that our approach reduces the complexity of the reductions in our programs to their optimal complexity for all of the 11 programs evaluated. In each 10 out of the 11 programs, the complexity improves by a (multiplicative) factor
of at least $N$, where $N$ is the size of the input data \(^1\). This is significant because for typical real application inputs of the programs in consideration, $N$ is in the range of $10^4$ to $10^6$ – a factor that subsumes other potential constant factor improvements. We also confirm this significance by showing that the speedups in wall-clock time ranges from 1.1x to over $10^7$x, with a median of 43x. We also outline the limits of the optimality of our approach, noting that our technique is not optimal if a reduction operator lacks an inverse operation.

In summary, multiple-statement reduction is a key ingredient of probabilistic inference algorithms, which are driving an emerging class of new programming languages and systems [Bingham et al. 2018; Cusumano-Towner et al. 2019; Daniel Huang 2017; Gelman et al. 2015; Goodman and Stuhlmüller 2014; Mansingkha et al. 2018; Narayanan et al. 2016; Tran et al. 2017] designed to streamline science and enable new applications. Optimizing these algorithms has historically either been done by hand or has been baked in as a domain/algorithmic-specification optimization for a single problem model [Holmes et al. 2012; Liu 1994]. To the best of our knowledge, our results are the first to identify and formulate multiple-statement reductions as a general program pattern, detail their challenges, and propose a technique to optimize their performance.

Road Map. In Section 2, We illustrate a heuristic algorithm to address the multiple-statement reduction described in Section 1. In addition, to further motivate the problem in the context of existing well-known algorithms, we present another motivating example which will be used for evaluation later in the paper. In Sections 3 and 4, we review backgrounds on polyhedral model and SR, respectively. In Section 5 we formalize our problem as a integer bilinear program. In Section 6 we introduce the proposed heuristic algorithm. In Sections 7 and 8 we discuss the implementation of our proposed algorithm and its evaluation. In Sections 9 and 10 we summarize some related work with concluding remarks.

2 EXAMPLE

In this section we will give two examples. In Section 2.1, we walk through our approach with the first example. In Section 2.2, we use a practical application example to further motivate the importance of the multiple-statement reductions problem.

2.1 Walk Through

In this section, we use the example of Equation (4) to 1) illustrate the steps of ST applications given reuse directions, 2) illustrate the invalid ST application that leads to dependency cycles, and compare it to the valid ST application, using the algorithm proposed in [Gautam and Rajopadhye 2006], and 3) describe the mechanism of our proposed heuristic algorithm, following the intuition we get from the comparison in 2).

**Naive Prefix Sum.** For ease of comparison and better visualization, we present the input in Equation (4) with Figure 1, a visual, polyhedral interpretation of the naive prefix sum program in Equation (4). In Figure 1, the top polyhedron with red dots represents the iteration domain of the reduction statement, $B[i] += A[j]$, with each red

\(^1\)For programs we consider, for example, this is usually the number of data points or the number of words of a text corpus. We include a more detailed review of input sizes for each benchmark in Section 8.3.2
dot denoting an iteration instance of the statement. The bottom polyhedron with blue squares represents the iteration domain for the statement \( A[i + 1] = f(B[i]) \). The middle polyhedron with orange diamonds is an additional polyhedron that our technique inserts into the program’s polyhedral representation to denote the completion of each reduction \( B[i] \).

**Data Dependencies.** Each arrow in Figure 1 represents a data dependence between iteration instances. An arrow from iteration instance \( a \) to instance \( b \) represents a data dependence from \( a \) to \( b \). The implication is that \( a \) needs to execute before \( b \).

There are three sources of data dependencies:

- **Reduction.** Each point in the middle polyhedron depends on all the points in the respective column of the top polyhedron. These dependencies are those of the reduction.

- **Use.** Each point in the bottom polyhedron depends on the point in the corresponding column of the middle polyhedron. These dependencies are those from the use of the reduction results.

- **Update.** Points in each row of the top polyhedron depend on the point in the bottom polyhedron that is one to the left of the leftmost point of the row. These dependencies are those induced by the update to \( A[i + 1] \) in Equation (4b) and use by Equation (4a).

**Incorrect Optimization.** The two diagrams presented in Figure 2 illustrate an incorrect application of ST using [Gautam and Rajopadhye 2006], which ignores the dependencies due to multiple-statement reduction.

Instead of using the correct reuse vector \([1, 0]^T\), this application uses the vector \([-1, 0]^T\). This vector maps iteration instances \([i, j]\) to instances \([i - 1, j]\). The organization of the diagram Figure 2a shows a mapping from the red shaded polyhedron the green shaded polyhedron, with each solid blue arrow represents the mapping between instances of the corresponding polyhedron. The green polygons outlined in red circles are the intersection of the two polyhedrons. Note that because the reuse vector has property that the evaluation of the reduction body \( A[j]\) are the same for any two points in the same row, the evaluation of a reduction over any column \( \text{col} \) in this intersection triangle must have the same value as the evaluation of a reduction over the column to the left of \( \text{col} \). Therefore the intersection part of the domain will be eliminated by ST, by reusing previously computed reductions (i.e. compute \( B[i] \) from \( B[i - 1] \) by incrementalizing using points not in the intersection). Figure 2b shows the pruned digram from Figure 2a. The top polyhedron now has the red circles at the rightmost column and the green polygon dots along the hypotenuse of the shifted domain. Note that the hypotenuse is restricted to the domain of projected domain of the reduction and does not include the point \([i, j] = [-1, 0]\).

Point \( d \) and the red circles column in the top polyhedron in Figure 2b correspond to the reduction that initializes \( B[N-1] \) in Equation (3a). All points in the middle polyhedron except \( d \) then correspond to Equation (3b), i.e., each \( B[j] \) is computed by subtracting the successor point \( B[i+1] \) by \( A[i] \). Dependencies in Figure 2b are preserved from Figure 2a, with the newly introduced dependency along the reuse vector in the middle polyhedron, which is represented by solid orange arrows pointing to the left. However, as mentioned in Section 1, Figure 2b’s dependencies form cycles; for instance, points \( a, b, c, d \) forms a cycle. Therefore, the transformed program in Figure 2b does not have a valid schedule, and consequently the application of ST along the reuse vector with mapping \([i, j] \rightarrow [i - 1, j]\) produces an incorrect optimization.

**Correct Optimization.** Figure 3 presents two diagrams, corresponding to the two steps to correctly applying ST in [Gautam and Rajopadhye 2006], respectively. Figure 3a illustrates the first step of the algorithm, where the algorithm chooses a reuse vector and shifts the reduction statement’s iteration domain along the vector. That is, Figure 3a illustrates the shift along the reuse vector, \([1, 0]^T\), which maps iteration instances \([i, j]\) to instances \([i + 1, j]\). As shown in this figure, this
corresponds to the mapping of top polyhedron, colored in red, to its shifted counterpart, colored in green. Each solid blue arrow represents the mapping from an instance in the red polyhedron to its counterpart in the green polyhedron. The green polygons outlined in red circles are again the points in the intersection of the two polyhedrons, which, same as the previous example of incorrect ST application, will be eliminated by ST.

Figure 3b corresponds to the resulting polyhedron and dependence structure after ST eliminates redundant computations, by applying the correct ST with reuse vector $[1, 0]^T$ to Equation (4) (Equations (2a), (2b) and (4b)). Each instance in the intersection of the two polyhedrons has been eliminated, along with its induced dependencies. The middle polyhedron also has new dependence edges: an edge has been added between reduction instances along the direction of the reuse vector. This polyhedron denotes the iteration domain of Line 3 in Listing 3. Each new dependence edge therefore reflects that each $B[i]$ is computed from $B[i - 1]$.

**Heuristic for choosing a valid direction.** As we have seen from the previous illustration, it is important to choose a valid reuse vector with multiple-statement reductions. In this work, we propose a heuristic algorithm for choosing a valid reuse vector. Notably, one key difference between Figures 2 and 3 is the dependencies drawn on the middle polyhedron. Specifically, in the middle polyhedron of Figure 3, the drawn dependencies on $B[i]$ respects the scheduled computation order...
of \mathcal{B}[i] of the original program in Figure 1, whereas that of Figure 2 disobeys that scheduled order. This observation has inspired the heuristic algorithm that always chooses the reuse vector that is consistent with the scheduled computation order of the LHS of the reduction. We show that the reuse vector chosen with this algorithm is 1) always sound, and 2) guarantees optimality if each reduction operator in the target program has an inverse.

### 2.2 Simplifying Multiple-Statement Reductions in Practice

As we later show in Section 8 by studying a variety of benchmarks, multiple-statement reductions commonly appear in the specifications of many problems and algorithms across statistics, artificial intelligence (AI), and machine learning (ML) with applications to computer vision, physics, and medicine. However, the common practice is to develop these algorithms by hand. Therefore, our technique offers the opportunity to automatically translate a specification to an efficient implementation. In this section, we illustrate our technique on a fundamental computation used across statistics, AI, and ML.

**Specification and Implementation.** Consider the following specification of *Gibbs Sampling* [Geman and Geman 1984] on a two-cluster Gaussian Mixture Model [see for example, Murphy 2012] (GS-2GM). This computation is designed to cluster data points such that similar data points, alternatively *observations*, are assigned to the same cluster. The input to GS-2GM is a float array \( \text{Obs} \) that represents the observations. The two-cluster Gaussian Mixture Model (GMM) assumes that each single observation belongs to one of the two clusters, and that each cluster follows a Gaussian distribution. The goal of GS-2GM is to sample the array \( Z \) that represents the cluster membership of each of the given observations, following the desired GMM distribution. It achieves this goal by iteratively taking in an old cluster assignment for each observation in turn, and samples a new one by updating the assignments of the remaining observations. This process will produce a stream of sample of \( Z \)s that approach the true distribution of \( Z \). The mathematical specification of GS-2GM is given in Equation (5).

\[
P_o(z, i) \equiv P(o_{\text{obs}} | o_{\text{obs}} \backslash i, Z_{\backslash i}, Z_i = z) \tag{5a}
\]
\[
C_{zi} = \sum_{\forall j \text{ s.t. } j \neq i \land Z_j = z} 1, \forall z, i \tag{5b}
\]
\[
S_{zi} = \sum_{\forall j \text{ s.t. } j \neq i \land Z_j = z} o_{\text{obs}_j}, \forall z, i \tag{5c}
\]
\[
P_o(z, i) = N(\frac{S_{zi}}{C_{zi}} (1 + C_{zi})^{-1} + 1) \tag{5d}
\]
\[
P(Z_i = 0 | Z_{\backslash i}, o_{\text{obs}}) = \frac{P_o(0, 0)}{P_o(0, 0) + P_o(1, 1)} \tag{5e}
\]
\[
Z_i \sim P(Z_i | Z_{\backslash i}, o_{\text{obs}}), \forall i \in [1, N] \tag{5f}
\]

In Equation (5), we use the notation \( \backslash \) i to denote the set \{ \( j \mid j \in [1, N], j \neq i \) \}. Equation (5a) defines the function \( P_o(z, i) \) as an abbreviation of the distribution of \( o_{\text{obs}_i} \) given all values of \( o_{\text{obs}} \) except \( o_{\text{obs}_i} \) and all current assignments of \( Z \)s except \( Z_i \) to \( z \). In Equations (5b) and (5c), \( C_{zi} \) and \( S_{zi} \) represent the counts and sums, respectively, of all the observations except the one with index \( i \), for which the current old assignment of cluster membership is 0 (and similarly for \( C_{zi} \) and \( S_{zi} \), with membership of 1). Then a distribution \( P(Z_i | Z_{\backslash i}, o_{\text{obs}}) \) is defined by Equations (5d) and (5e). In this example, the distribution is simply tuple of two floats representing the weights of assigning \( Z_i \) to cluster zero or one, respectively. Note that the exact computation required to produce this tuple is not important for understanding the optimization problem. The key information is that they are produced by a deterministic pure function that depends only on the counts and sums defined above. Lastly, Equation (5f) samples each \( Z_i \) in order from this distribution. Listing 4 gives an efficient
implementation of the above mathematical specification – notice that Listing 4 computes the counts and sums incrementally, instead of forming the full reductions of Equations (5b) and (5c). Deriving Listing 4 from Equation (5) requires manually solving the multiple-statement reductions problem which is tedious and error-prone.

**Our Approach.** Given an array-based representation of Equation (5), our approach automatically produces Listing 4. For conciseness of presentation, we consider the variable $S_{z_i}$ with fixed $z = 0$ as an example. In this case Equation (5c) can be rewritten as sum of two variables $S_{0i} = S_{0L}[i] + S_{0R}[i]$, where $S_{0L}$, $S_{0R}$ are given by Equations (6a) and (6b), respectively. The step of rewriting in terms of $S_{0L}$ and $S_{0R}$ is standard in polyhedral model compilation: the original domain with constraint $j \neq i$ is non-convex and it is standard to break it into two convex polyhedrons with constraints $j < i$ and $j > i$. Further, we make the non-affine constraint $Z_j = z$ into a simple if-then-else expression guarding the reduction’s body – this is standard approach and same as the one proposed by Benabderrahmane et al. [2010] to model non-affine constraints as control predicates.

\[
S_{0L}[i] = \sum_{j=0}^{j<i} (Z[j] == 0 ? \text{Obs}[j] : 0) \tag{6a}
\]

\[
S_{0R}[i] = \sum_{j=i+1}^{j<N} (Z[j] == 0 ? \text{Obs}[j] : 0) \tag{6b}
\]

\[
\vdots \quad \text{Other equations...}
\]

\[
Z'[i] = \text{sample}(S_{0L}[i] + S_{0R}[i], ...) \tag{6c}
\]
Equations (6a) and (6c) exactly correspond to Equations (4a) and (4b), respectively, since they have the same data flow dependencies. Thus the technique walked through in Section 2.1 also applies to Equations (6a) and (6c) to produce a specification with efficient complexity. Further, our technique is general in that it handles any multiple-statement reduction, including Equation (6b) with constraints $i + 1 \leq j < N$, where the constraints are the reverse of the constraints in Equation (6a). Lastly, the same analysis can be applied to all cases of $C_{ij}$ and $S_{ij}$ with $z = 0$ or $z = 1$. The analyses in total produces eight intermediate variables, namely $C_{0L}, C_{1L}, C_{0R}, C_{1R}, S_{0L}, S_{1L}, S_{0R}, S_{1R}$, which produce Listing 4 by applying our technique and compiling to executable code.

**Results.** Our evaluation shows that our technique produces an optimal complexity algorithm for Gibbs Sampling on the Gaussian Mixture Model, matching that of a manually developed implementation, and yielding a 7.4x performance improvement over a naive, unoptimized implementation. These results demonstrate the opportunity to automatically compile high-level specifications that include multiple-statement reductions to efficient implementations.

### 3 BACKGROUND: POLYHEDRAL MODEL

In this section, we review terminologies from the polyhedral model that we use in this work.

#### 3.1 Polyhedral Set Representation

We use the following definition and notation for a polyhedral set; the notation is consistent with the Integer Set Library (ISL) [Verdoolaege 2010]’s notation.

**Definition 3.1 (System of affine inequalities).** A system of affine inequalities is defined as $A \cdot \overline{x} \geq 0$: $A$ is an $m \times (n + 1)$ constant integer matrix and $\overline{x}$ is length-$n$ vector of integer unknowns.

**Remark 1.** We may also express a system of affine inequalities by conjunction of simple affine inequalities. For example, the system $\begin{bmatrix} 1 & 0 & 0 \\ 1 & -1 & 1 \end{bmatrix} \cdot \overline{x} \geq 0$ is equivalent to $(x \geq 0) \land (x \geq N - 1)$ — or simply the short hand $0 \leq x < N$. A simple equality $x = 0$ is short hand for the conjunction of two inequalities $(x \geq 0) \land (\neg x \geq 0)$.

**Definition 3.2 (Polyhedral set).** A polyhedral set $\mathcal{P}$, defined as $[\vec{p}] \rightarrow \{[\overline{x}] : A \cdot [\overline{x}, \vec{p}, 1]^T \geq 0\}$, contains a tuple of parameters $[\vec{p}]$, a tuple template $[\overline{x}]$ and a system of affine inequalities $A \cdot [\overline{x}, \vec{p}, 1]^T \geq \vec{0}$, where $A$ is a matrix of coefficients. We say $[\vec{p}] \rightarrow \{[\overline{x}]\}$ is the space of $\mathcal{P}$.

For example, $[N] \rightarrow \{[i] : 0 \leq i < N\}$ denotes the set of integers from 0 to $N - 1$. The space of this set is $[N] \rightarrow \{[i]\}$.

**Definition 3.3 (Polyhedral relation).** A polyhedral relation $[\vec{p}] \rightarrow \{[\overline{x}_1] \rightarrow [\overline{x}_2] : A \cdot [\overline{x}_1, \overline{x}_2, \vec{p}, 1]^T \geq \vec{0}\}$ contains a tuple of parameters $[\vec{p}]$, tuple templates $[\overline{x}_1], [\overline{x}_2]$ and a system of affine inequalities $A \cdot [\overline{x}_1, \overline{x}_2, \vec{p}, 1]^T \geq \vec{0}$.

For example, $[N] \rightarrow \{[i, j] \rightarrow [i + 1, j] : 0 \leq i < N, 0 \leq j < N\}$ denotes the relation that maps every integer tuple $[i, j]$ to $[i + 1, j]$ within an $N$-by-$N$ grid. Figure 4 visualizes this relation with $N = 5$: the blue arrows map points corresponding to integer tuples to their right successors.

---

2 Although Equation (6c) contains sample that is stochastic and Equation (4b) contains $f$ that is deterministic, they still have the same data flow dependencies.
Semantically, the polyhedral set provides an intentional description for a set of tuples, templated by \( [\vec{x}] \), so that all tuples in the set satisfy the system of affine inequalities. The set is optionally parametric in \( [\vec{p}] \), if \( [\vec{p}] \) is not empty.

Similarly, a polyhedral relation describes a set of binary relations mapping from \( [\vec{x}_1] \) to \( [\vec{x}_2] \), for every \( [\vec{x}_1]-[\vec{x}_2] \) pair that satisfies the system of affine inequalities; a polyhedral relation can also be parametric in \( [\vec{p}] \). For aesthetic reasons, we omit the parameter \( [\vec{p}] \) when it is clear from the context which identifiers are parameters.

**Definition 3.4 (Face of polyhedral set).** Let the polyhedral set \( \mathcal{P} = [\vec{p}] \rightarrow \{[\vec{x}] : A \cdot [\vec{x}, \vec{p}, 1] \geq \vec{0} \} \). Let \( M_i \) be the \( i \)-th row of matrix \( M \). A face of \( \mathcal{P} \) is defined as \( \mathcal{F} = \mathcal{P} \cap \mathcal{B} \) where \( \mathcal{B} = [\vec{p}] \rightarrow \{[\vec{x}] : B \cdot [\vec{x}, \vec{p}, 1] = \vec{0} \} \) and \( \forall i \exists j, A_i = B_j \).

In words, a face of \( \mathcal{P} \) is \( \mathcal{P} \) with a subset of (potentially empty or all) inequalities of \( \mathcal{P} \) changed to equality.

### 3.2 Polyhedral Representation of a Program

The polyhedral model represents a program by a set of statements, and for each statement, an associating polyhedral set known as the statement’s domain. Each point in a polyhedral set correspond to one concrete execution instance of the statement.

#### 3.2.1 IR Syntax.** Following the formalization by the original SR work [Gautam and Rajopadhye 2006; Yuki et al. 2013], we use an equation-based representation of program in this work, presented in grammar by Listing 5. We explain each component in turn:

- **<prog>** a program consists of multiple statements.
- **<stmt>** a statement is left hand side (i.e. \( \text{LHS[<afflist>]} \)), middle assignment operator (i.e. either \( = \) or \( \oplus = \) ), a right hand side expression (i.e. \( <\text{expr}> \)), and its domain (i.e. \( \mathcal{P} \)). A statement is a normal assignment statement when the middle assignment operator is plain \( = \); a statement is a reduction when the middle assignment operator is \( \oplus = \).
- **<expr>** an expression is either an unary or binary operator applied on expression(s), an array reference (i.e. \( \text{ARR[<afflist>]} \)), or a constant.
- **<aff>** an affine expression is a kind of expression that applies affine transformation to variables and produces a scalar. It references only variables in \( \vec{x} \) or \( \vec{p} \), where \( [\vec{p}] \rightarrow \{[\vec{x}] \} \) is the space of \( \mathcal{P} \).
- **<afflist>** a list of affine expressions. Array references (i.e. \( \text{LHS[<afflist>]} \) and \( \text{ARR[<afflist>]} \)) must have indices that are affine expressions. An \( <\text{afflist}> \) of length \( n \) can be expressed mathematically as an affine transformation \( A \cdot [\vec{x}, \vec{p}, 1]^T \), where \( A \) is a constant \( n \times (|\vec{x}| + |\vec{p}| + 1) \) integer matrix and \( \vec{x}, \vec{p} \) defined same as those for \( <\text{aff}> \).
- **\( \mathcal{P} \)** a polyhedral set representing the statement’s domain. Since each point in the domain corresponds to one concrete execution instance of the statement, if \( \mathcal{P} = [\vec{p}] \rightarrow \{[t] : e \} \), then \( \vec{p} \) corresponds to the set of parameters of the program and \( t \) corresponds to the set of loop variables of the statement.
3.2.2 IR Semantics.

Access Relation. An access relation is a polyhedral relation mapping from the space of a statement’s domain to the space of an accessed array. An access relation can either be a write access relation (in case of LHS<afflist>), or a read access relation (in case of RHS<afflist>). Let ARR<afflist> be an array reference for a statement with space \([\vec{p}] \to \{[\vec{x}]\}\) and <afflist> expressed as \(A \cdot [\vec{x}, \vec{p}, 1]\), the access relation for this array access is \([\vec{p}] \to \{[\vec{x}] \to [\vec{y}] : A \cdot [\vec{x}, \vec{p}, 1] = [\vec{y}]\) in case of LHS<afflist>, or \(\{[\vec{x}] \to [\vec{y}] : A \cdot [\vec{x}, \vec{p}, 1] = [\vec{y}]\) in case of RHS<afflist>.

Reduction projection. If a statement is a reduction, we define the projection of the reduction proj as the write access relation of LHS array reference of the reduction.

SSA. Following [Gautam and Rajopadhye 2006], our IR requires the program to be in array static-single-assignment (Array SSA) form[Feautrier 1988]; that is, each array element is never written twice during program execution. To our IR, this means for each unique LHS array, and the statements \(S_0 \ldots S_k\) that writes to it, \(\bigcap_i W_{S_i} = \emptyset\), where \(W_{S_i}\) is the write access relation for \(S_i\).

Semantics. We use usual semantics from array languages [Yuki et al. 2013] for our IR. Specifically, a statement is evaluated under each point of its domain \(P\). An expression is evaluated under a point by substituting the free variables of the expression with the instantiated values of those variables under that point. For example, \(A[\vec{N} - i + 1]\) evaluates to the value of \(A[9]\) at point \([\vec{N}] \to \{[i] : N = 10 \land i = 2\}\). If the statement is a normal assignment, for each point in \(P\), the right hand side expression is evaluated and assigned to the left hand side array. If the statement is a reduction, for each point \(p \in P\) the right hand side expression is evaluated, and its value is accumulated into LHS at point \(p' = proj(p)\) using the operator \(\oplus\) where \(proj\) is the projection of the reduction as defined previously.

3.3 Polyhedral Model Scheduling

Scheduling is a step in polyhedral model where a scheduling function assigns each point in a statement’s domain a timestamp, denoting the order of all execution instances. This step is essential for multiple-statement programs because the timestamps are assigned to respect the inter-statement and intra-statement dependencies.

3.3.1 Scheduling Function.

Definition 3.5 (Schedule Timestamp). A schedule timestamp is an \(m\)-dimensional vector, where \(m\) is the upper bound on the dimension of the schedule. For two timestamps \(T_1\) and \(T_2\), \(T_1 < T_2\) (\(T_1\) happens before \(T_2\)) iff \(T_1[i] < T_2[i]\) where \(i\) is the first non-equal index between \(T_1\), \(T_2\).

A schedule \(\Theta\) for a program is a collection of scheduling functions, one for each statement. A scheduling function for a statement \(S\) is an affine transformation, represented by the matrix \(\Theta_S\), which maps statement \(S\)’s domain to its scheduling timestamp. For a statement \(S\) with domain in space \([\vec{p}] \to \{[\vec{x}]\}\), its \(m\) dimensional timestamp \(T_S\) is given by the \(m \times (|\vec{x}| + |\vec{p}| + 1)\) scheduling function \(\Theta_S\):

\[
T_S = \Theta_S \cdot \begin{bmatrix} \vec{x} \\ \vec{p} \\ 1 \end{bmatrix} = \begin{bmatrix} \theta_{1,1} & \cdots & \theta_{1,|\vec{x}|+|\vec{p}|+1} \\ \vdots & \ddots & \vdots \\ \theta_{m,1} & \cdots & \theta_{m,|\vec{x}|+|\vec{p}|+1} \end{bmatrix} \cdot \begin{bmatrix} \vec{x} \\ \vec{p} \\ 1 \end{bmatrix}
\]

(7)

3.3.2 Dependence Relation. Any two statements, \(S, T\), must satisfy a dependence relation represented by a polyhedral relation \(D_{S,T} = [\vec{p}] \to \{[\vec{x}_1] \to [\vec{x}_2] : D_{S,T} : [\vec{x}_1, \vec{x}_2, \vec{p}, 1]^T \geq \vec{0}\}\), and \(D_{S,T}\) is the dependency matrix. The dependence relation \(D_{S,T}\) describes the happens before relation between iterations of \(S\) and \(T\). For a pair of statements \(S, T\), let \(S\) writes to LHS and \(T\)’s RHS expression reads
elements of LHS. The dependence relation $D_{S,T}$ is equal to $R^{-1} \circ W$, where $R$, $W$ are the read and write access relations for LHS of the two statements respectively, $R^{-1}$ denotes the inverse of the polyhedral relation $R$, and $\circ$ denotes composition. Previous work [Collard et al. 1995; Verdoolaege et al. 2013] and textbook [Verdoolaege 2016] contain detailed introductions to dependence analysis techniques, which we refer to the reader for a deeper exposure.

### 3.3.3 ILP formulation of scheduling

The task of scheduling a program in polyhedral model is to find a schedule $\Theta$ for the program such that the schedule timestamps for all statements instances satisfy the dependence relations of the program. Pouchet et al. [2011] formalized the scheduling problem for obtaining $m$-dimensional schedule as the following convex problem:

\[
\forall D_{S,T}, \forall k, \delta_k^{D_{S,T}} \in \{0,1\} \tag{8a}
\]

\[
\forall D_{S,T}, \sum_{k=1}^{m} \delta_k^{D_{S,T}} = 1 \tag{8b}
\]

\[
\forall D_{S,T}, \forall k \in [1,m], \forall [\vec{x}_S, \vec{x}_T, \vec{p}] \in D_{S,T} \tag{8c}
\]

\[
\Theta^k_S \cdot \begin{bmatrix} \vec{x}_S \\ \vec{p} \\ 1 \end{bmatrix} - \Theta^k_T \cdot \begin{bmatrix} \vec{x}_T \\ \vec{p} \\ 1 \end{bmatrix} \geq - \sum_{i=1}^{k-1} \delta_i^{D_{S,T}} (K \vec{p} + K) + \delta_k^{D_{S,T}} \tag{8d}
\]

In words, the formulation creates a binary variable $\delta_k^{D_{S,T}}$ for each $k \in [1,m]$ dimensions, and each pair of dependence relation in the program. The binary variable is used to model entry-by-entry comparison of an $m$ dimensional timestamp. Constraint c) finally encodes that the schedule function $\Theta_S$ and $\Theta_T$ must satisfy that $\vec{x}_S$ is scheduled before $\vec{x}_T$, if the dependence $\vec{x}_S \to \vec{y}_T$ exists — that is, $[\vec{x}_S, \vec{x}_T, \vec{p}]^T \in D_{S,T}$. The variable $K$ is a known constant obtainable from the original program, and is an upper bound modeling technique to make the problem convex. Pouchet et al. [2011] shows that this problem is equivalent to an ILP thanks to Farkas’ Lemma [Schrijver 1986], solving which produces the desired schedule coefficients $\Theta$ in Section 3.3.1.

## 4 BACKGROUND: THE SIMPLIFYING REDUCTIONS FRAMEWORK

Previous work [Gautam and Rajopadhye 2006] introduced a core transformation called the simplification transformation (ST) that can potentially transform a single statement specified in Listing 5 to lower its complexity, along with a set of enabling transformations: reduction decomposition, same operator transformation, distributivity transformation and higher-order operator transformation. For the core transformation, we will use an example from Section 1 to illustrate the transformation. For the enabling transformations, we will include a brief description for each transformation. Finally, [Gautam and Rajopadhye 2006] combine all the transformations to provide a dynamic programming algorithm to efficiently choose from an infinite set of configurations and orders for the transformations, a sequence of transformations that lead to optimal complexity reduction.

### 4.1 Simplification Transformation

Here we use take the example from Section 2.1 to illustrate the core simplification transformation that reduces complexity of a reduction. The full specification of ST can be found in Appendix B and Gautam and Rajopadhye [2006].

Listing 6 illustrates the example of applying ST (Section 3.2) to Equation (4) and producing the optimized version (Equations (2a), (2b) and (4b)) in our IR. As we mentioned before, core ST operates on single statement only, and produces correct result for multiple statements if a correct reuse vector is given.
Simplifying Multiple-Statement Reductions with the Polyhedral Model

Listing 6. ST in the polyhedral IR for the example in Section 1 (Equation (4)), given the reuse vector \([1, 0]^T\)

Original Reduction. Listing 6 presents the reduction in Equation (4a) in the polyhedral IR as the statement S1Add, which provides the contents of BTmpAdd. Specifically, S1Add is a polyhedron over the full space of \(i\) that sets BTmpAdd\([i]\) to equal \(A[i]\). TAddOn1y next initializes BTmp\([0]\) (as on Line 1 of Listing 3) and S1AddReuse incrementally computes the remaining values of BTmp (as on Line 3 of Listing 3).

Algorithm (Reuse Vector). To identify this optimization opportunity and generate the optimized code, the Simplification Transformation identifies a reuse vector by which shifting the original, unoptimized polyhedron \(P\) makes plain that consecutive iterations of the polyhedral overlap and can therefore be incrementalized.

Consider the reuse vector \(\vec{r} = [1, 0]^T\), that shifts all points \([i, j]\) to \([i + 1, j]\); \(\vec{r}\) can also be represented by the polyhedral relation \([i, j] \rightarrow [i + 1, j] : \forall i, j\). The arrows in Figure 5 visualizes \(\vec{r}\) over the domain of the original reduction (red dots in the shaded red triangle).

Given this reuse vector, ST performs the following steps:

- **Shift.** The transformation first shifts T’s polyhedron along the direction of the reuse vector, transforming \([i, j] : 0 \leq i < N \land 0 \leq j \leq i\) (red dots in Figure 5) into \([i, j] : 1 \leq i < N + 1 \land 0 \leq j \leq i - 1\) (green hexagon points in the shaded green triangle in Figure 5).

- **Intersect.** The transformation next computes the intersection of the shifted polyhedron with its original polyhedral, yielding \([i, j] : 1 \leq i < N \land 0 \leq j \leq i - 1\) (overlapped points in the shaded purple triangle in Figure 5). This polyhedral denotes the subset of points of the original domain \(P\), whose value can be reused from the predecessor points as indicated by the reuse vector.

- **Project.** Finally, the transformation projects the result onto the space of polyhedron that represents the indices of left hand side array \(S\). Concretely, the transformation applies the projection represented by the polyhedral relation \([i, j] \rightarrow [i] : \forall i, j\), yielding the polyhedron \([i] : 1 \leq i < N\).

This final polyhedron is exactly the domain of elements of \(S\) that exhibits reuse along the reuse vector \(\vec{r}\). For this example, this means that instead of computing \(S[i]\) with the original accumulation \(S[i] := A[i]\), the transformation computes \(S[i] = S[i - 1] + A[i]\) on the final polyhedral domain \([i] : 1 \leq i < N\).
Completion. The polyhedral $\{ [i] : 1 \leq i < N \}$ does not cover the full domain of the original reduction. Specifically, it is missing $S[i]$ on the domain $\{ [i] : i = 0 \}$ – that is, exactly when $i = 0$. The value of $S[0]$ should be equal to $A[0]$. Therefore, we should expect two IR statements: one statement for initializing $S[0]$ to $A[0]$, and one statement incrementing while reopening a previous value of $S[i - 1]$.

The transformed code in Listing 6 captures this intuition: $T$-add-only for initialization and $T$-add-reuse for incrementing and reusing; however, instead of directly using $A[j]$, the transformation uses an auxiliary array $S\_ADD$, which is useful for generalizing to cases where initialization or incrementization requires more than one value from the right hand side expression (i.e. $A$).

4.2 Configuration of Simplification Transformation

A fully automated optimizing compiler should automatically identify a reuse direction $\vec{r}$ and apply ST. There are several considerations when comes to choosing $\vec{r}$.

- **Complexity**: performing ST along $\vec{r}$ reduces the complexity of the computation.
- **Inverse**: if the inverse of $\oplus$ does not exist, then performing ST along $\vec{r}$ will not require an inverse operator.
- **Sharing**: RHS expression presents sharing (defined below) along $\vec{r}$.

Each requirement prescribes a set, potentially an infinite set, of vectors $\vec{r}$, which we explain below:

**Complexity.** We require that applying ST along reuse vector $\vec{r}$ reduces the complexity of the program. The complexity of a program will not increase after applying ST for any $\vec{r}$; however, the complexity can stay the same if $\vec{r}$ is chosen along a direction where the original polyhedral domain $\mathcal{P}$ has constant thickness – that is, the extent of $\mathcal{P}$ is bounded by some constant not parameterized by the input parameters of the program. For example, consider an extreme case of the prefix sum example (Listing 1, Listing 2) but with the input parameter $N$ fixed to some constant – say $N = 4$. The complexities before and after ST will be the same – $O(1)$ – since both programs will perform a fixed number of computations. For a statement $S$ with domain $\mathcal{P}$, we use $L(\mathcal{P})$ to denote the set of vectors $\vec{r}$ that satisfies the complexity condition.

**Inverse.** If $\oplus$ does not have an inverse, we require that applying ST along a vector $\vec{r}$ will not introduce statements that requires the inverse operator of $\oplus$. For example, if $\oplus$ is min() or max(), it does not have an inverse; in such cases, [Gautam and Rajopadhye 2006] introduces the concept of Boundary Constraints – which in short is the set of constraints of the domain $\mathcal{P}$ that are orthogonal to the projection $\text{proj}$ – and require that $\vec{r}$ must be pointing out of (instead of pointing into) the boundaries of $\mathcal{P}$ corresponding to the Boundary Constraints. For a statement $S$, we use $I(S)$ to denote the set of vectors $\vec{r}$ that satisfies the inverse condition.

**Sharing.** We require that applying ST along a vector $\vec{r}$ where the right hand side expression of the considered reduction presents sharing along $\vec{r}$. For example in Section 4.1, we analyzed that the prefix sum example has sharing along direction of $i$. [Gautam and Rajopadhye 2006] introduced an algorithm to determine the share space, the space formed by all reuse directions, given an equationally specified reduction. For a statement $S$, we use $S(S)$ to denote the set of vectors $\vec{r}$ that satisfies the sharing condition.

In general, for a statement $S$, denote its domain as $S\_domain$; we would like to find the intersection $\mathcal{R}(S) = L(S\_domain) \cap I(S) \cap S(S)$, so that any $\vec{r} \in \mathcal{R}(S)$ is a valid reuse vector to perform ST.
4.3 Recursive ST

Notice in Listing 6 that statement Tadd still contains a reduction. Although for this example Tadd does not have further ST opportunities, in general, the residual reduction might still have available ST opportunities so that ST can be applied recursively to all introduced reductions.3

5 MULTIPLE-STATEMENT SR PROBLEM

In this section, we state the Multiple-Statement Simplifying Reduction (MSSR) problem. In particular, we focus on the core of the Simplifying Reduction approach – the Simplification Transformation in Section 4.1 – and do not consider the Simplifying Reduction framework’s additional enabling transformations. These transformations increase available simplification opportunities; we briefly touch on enabling transformations in Section 5.3.

5.1 Problem Statement

Ideally we would formulate the Multiple Statement Simplifying Reduction (MSSR) as Equation (9).

\[
\begin{align*}
\text{minimize} & \quad \text{complexity}(\text{prog}') & \text{(9a)} \\
\text{subject to} & \\
\text{prog}^1 & = \text{prog}, \quad \text{prog}' = \text{prog}^n & \text{(9b)} \\
\forall i \in [2, n]: & \quad \text{prog}^i = \text{ST}_{S_i, \vec{r}_i}(\text{prog}^{i-1}) & \text{(9c)} \\
\vec{r}_i & \in \mathcal{L}(S_i, \text{domain}) \cap \mathcal{I}(S_i) \cap \mathcal{S}(S_i) & \text{(9d)} \\
\exists & \text{ schedule } \Theta \text{ of } \text{prog}' , & \text{(9e)} \\
\text{s.t. } & \Theta \text{ satisfies } \text{dependence}(\text{prog}') & \text{(9f)} \\
\text{given } & \text{prog, dependence}(\text{prog}) & \text{(9g)} \\
\text{variables } & S_1, S_2, \ldots, S_n, \vec{r}_1, \vec{r}_2, \ldots, \vec{r}_n & \text{(9h)}
\end{align*}
\]

This states that given a program prog, and all pairwise dependencies between those statements, dependence (prog), apply a sequence of n ST transformations, ST_{S_i, \vec{r}_i} that minimizes the complexity of the resulting program, prog'. Here we use ST_{S_i, \vec{r}}(prog) to denote an ST that is applied on a statement S in prog along the reuse vector \vec{r}. Further, Equation (9d) requires each \vec{r}_i to satisfy the constraints (i.e. complexity, inverse and sharing, denoted by \mathcal{L}(.), \mathcal{I}(.), \mathcal{S}(.) respectively) as stated in Section 4.2.

Unfortunately, Equation (9) has three issues: 1) it is not a well-defined formulation due to the unknown n 2) it has infinite space for \vec{r}_i 3) it has impractically large space for S_i.

First, it is not a well-defined formulation: to define its variables, the problem relies on an oracle to produce n, the total number of ST applications – even though there is no readily apparent bound on that number. To elaborate, each ST application removes one statement, and introduces zero to two reductions that are potentially applicable for further ST applications – thus one needs justification that recursively applying ST always terminates in order to bound n.

Secondly, even if we assume that n is given and bounded, the formulation does not readily translate to an executable algorithm. Specifically, enumeratively searching all possible \vec{r}_i combinations is not feasible: each \vec{r}_i alone is chosen from an infinite set of vectors, and the entire search space is also infinite; therefore the search space of \vec{r}_i's is impossible to navigate with enumerative search.

Thirdly, also assuming n is given and bounded, the program relies on a sequence of S_i to specify on which statement in prog' to perform ST. Although, unlike the case of \vec{r}_i, the number of choices for each S_i is finitely bounded (i.e. by the number of ST-applicable reductions in the program), the combinations of all possible (S_1, ... S_n) has at least |S_i|! possibilities: assuming the best case scenario where each ST applications removes one reduction and introduces zero reductions that are

3In general, ST can also introduce more than one reduction; we include a full description of ST in Appendix B.
potentially applicable for further ST applications, which implies the $i$th ST application has $n - i + 1$ remaining alternative choices of $S_i$ (i.e. $|S_i| = n - i + 1$). Therefore the search space of $S_i$ is also not practical to navigate with enumerative search.

We will resolve these issues with a correct formalization in the rest of Section 5. Specifically, we show, for a program, a one-to-one correspondence between all its potential ST applications and all faces of its reductions’ domains. This correspondence resolves the first issue by bounding the number of ST applications to the number of faces of the program. This correspondence also allows a construction of an Integer Bilinear Programming (IBP) formulation to MSSR, which avoids the explicit enumerative search in the second and third issue.

5.1.1 Per-face ST application. We first make the following observation of ST on a single statement $S$ with domain $P$: if we apply ST on $S$, we can then recursively apply ST on the newly introduced reductions, as in Section 4.3, and this is exactly the root problem of the incorrect formulation Equation (9): this recursion appears non-terminating. We will solve this issue by stating and proving Lemma 5.2 — to this end, we first recall Lemma 5.1 from [Gautam and Rajopadhye 2006] that we will use in our proof. We then state Lemma 5.2 and give a proof.

**Lemma 5.1 (Local Face Correspondance [Gautam and Rajopadhye 2006, Theorem 3]).** Let $P'$ be the translation of an $n$-dimensional $P$ along $\hat{r}$, then $P - P' = \cup P_i$, and there exists a one-to-one map from $i$ to faces of $P$ such that each $P_i$ corresponds uniquely to a $(n - 1)$-dimensional face of $P$.

**Lemma 5.2 (Global Face Correspondance).** Each recursive application of ST is on a subset (a polyhedral set) of $P$, and all subsets correspond exactly one-to-one to all faces of $P$.

**Proof.** Given a statement $S$ with domain $P$, ST performs a shift of $P$ along a given reuse vector to $P'$; new reduction statements are introduced over domains $P - P'$ and $P' - P$. Note that these two domains are non-convex half shells around the original domain $P$, and together form a full shell around $P$. The two shells are both non-convex, however by Lemma 5.1, they decompose into convex polyhedral domains, each corresponding to a unique $(n - 1)$-dimensional face of the $n$-dimensional polyhedron $P$.

ST is applied recursively on these decomposed $(n - 1)$-dimensional faces and then on the sequence of $(n - i)$-dimensional faces until the recursion hits the vertices of $P$. The entire recursion is therefore a procedure that enumerates through all faces of a statement $S$’s full domain $P$ and assigns a reuse vector to each face.

With Lemma 5.2, the recursive ST application always terminates since the number of faces of $P$ is finite. Further, this introduces a per-face application view of ST — under this view, the algorithm first chooses a reuse vector for each face of $P$ up-front; it then uses the same recursive ST application starting at $P$ same as before; however, for each sub-domain’s ST application, it uses the reuse vector assigned for the face corresponding to that sub-domain. Lastly, note that the reuse vector assigned to each face is parallel to the face because the residual domain corresponding to the face already has constant thickness orthogonal to that face – therefore shifts not parallel to the face do not change complexity and disobey the rule stated for Complexity in Section 4.2.

5.2 Integer Bilinear Program Formulation

With the per-face application view of ST in Section 5.1.1, we are now ready to give the correct formulation of MSSR. The basic idea behind this formulation is to combine previous work on SR for a single statement [Gautam and Rajopadhye 2006], previous work on the integer linear program formulation of polyhedral model scheduling [Pouchet et al. 2008, 2007, 2011] and the per-face application view of ST presented in Section 5.1.1. We first revisit Equation (9) and give the correct
high level formulation as follows:

\begin{align}
\text{minimize} & \quad \text{complexity}(\text{prog}') \\
\text{subject to} & \quad \text{prog}' = (\text{ST}_{f_1, r_1} \circ \ldots \circ \text{ST}_{f_n, r_n})(\text{prog}) \\
& \quad \forall i \in [1, n]: \quad r_i \in \mathcal{L}(f_i) \cap \mathcal{I}(f_i, \text{stmt}) \cap \mathcal{S}(f_i, \text{stmt}) \\
& \quad \exists \text{ schedule } \Theta \text{ of } \text{prog}', \quad \Theta \text{ satisfies dependence}(\text{prog}') \\
& \quad \text{given } \text{prog}, \text{ dependence} (\text{prog}) \\
& \quad \text{variables } \vec{r}_1, \vec{r}_2, \ldots, \vec{r}_n
\end{align}

For Equation (10), \( \{f_1 \ldots f_n\} \) denotes the set of all faces of domains of all statements in \text{prog}. Following the per-face view of ST in Section 5.1.1, the function composition \( \text{ST}_{f_1, r_1} \circ \ldots \circ \text{ST}_{f_n, r_n} \) denotes applying per-face ST with the assigned reuse directions \( \vec{r}_1 \ldots \vec{r}_n \) (for all faces \( f_1 \ldots f_n \)). We use \( f \cdot \text{stmt} \) to denote a face \( f \)'s corresponding statement (i.e. the statement which has the domain \( f \)).

This high level formulation is similar to Equation (9), except that now 1) each reuse vector \( \vec{r}_i \) is in one-to-one correspondence with a face \( f_i \) — we thus have a bounded number of unknown variables for reuse vectors 2) the variables \( S_i \) are eliminated, as the new formulation uses the per-face ST view, instead of the recursive ST application view. Lastly, each reuse vector is still constrained to satisfy the validity constraints (i.e. Equation (10c)).

5.2.1 Variables. As we mention above, the unknown variables contains reuse vectors \( \vec{r}_1 \ldots \vec{r}_n \). Note that, moreover, the existential quantification over \( \Theta \) (Equation (10d)) implies that \( \Theta \) is also an unknown. As in Section 3.3.1, \( \Theta \) is a collection of scheduling functions \( \Theta_S \), one for each statement \( S \) in the final program \( \text{prog}' \); each \( \Theta_S \) is an \( m \times n \) dimension matrix of integer unknowns, where \( m \) is the schedule dimension, and \( n \) is one plus the sum of the number of dimension of \( S \cdot \text{domain} \) and the number of parameters of the program.

5.2.2 Constraints. Equation (10) contains two main categories of constraints: reuse constraints in Equation (10c) and dependency constraints in Equation (10f).

Reuse constraints. The reuse constraints enforce that each \( \vec{r}_i \) is chosen from \( \mathcal{L}(f_i) \cap \mathcal{I}(f_i, \text{stmt}) \cap \mathcal{S}(f_i, \text{stmt}) \) — this later set is a union of polyhedral sets computable from \( f_i \). Since it’s a union of polyhedral sets, we use disjunction to constrain \( \vec{r}_i \) to belong to one of the polyhedral sets. For each the polyhedral set, encoding that \( \vec{r}_i \) belongs to the polyhedral set is then just a simple affine inequality constraint.

Dependency constraints. The dependency constraints enforce that \( \Theta \) satisfies the dependency of \( \text{prog}' \). Specifically, it requires that for each pair of statements \( S \) and \( T \) that potentially occur in \( \text{prog}' \), their scheduling functions \( \Theta_S, \Theta_T \) satisfy the dependence relation \( D_{S,T} \). On the high level, we set up the constraints just the same as in Equation (8). However, the dependence matrix \( D_{S,T} \) now contains entries with (linear) terms with unknowns from \( \vec{r}_1 \ldots \vec{r}_n \). An informative argument for why \( D_{S,T} \) contains these unknown entries is: if we look from the recursive ST view, each application of ST introduces a reuse direction unknown \( \vec{r}_i \), and the algorithm recurses down to the residual reductions – for the next recursive application, we can think of it as taking in a program with both the original program’s parameters, and also the reuse vector unknowns introduced by the previous ST application. The residual reductions’ domains then have space extended by all \( \vec{r}_1 \ldots \vec{r}_n \).

5.2.3 Objective: complexity. Since we would like to minimize the overall complexity, we need to express our integer bilinear program’s objective as the complexity of the transformed program. We can compute complexity of each face by counting the cardinality of each face’s domain [Verdoolaege
The cardinality of a face is an Ehrhart polynomial [Ehrhardt 2009] in terms of the program parameters.

**Encoding.** If the program only has one parameter, then the degree of the polynomial is a natural choice of a scalar that represents the complexity of the program.

If the program has multiple parameters, then one needs to be careful about comparing complexities: it is necessary to be able to compare between $O(M^2N)$ and $O(MN^2)$ in order to minimize complexity. To this end, we assume that a total ordering is given for all possible polynomial terms of global parameters as a sequence of increasing scalars. For example, with two global parameters $M, N$, and maximum possible complexity $O(M^2N^2)$, a total ordering such as $O(1) < O(M) < O(N) < O(MN) < O(M^2N) < O(MN^2) < O(M^2N^2)$ is given, and integers $0...6$ are assigned to each big-O term in the previous sequence.

**Summing scalar encodings.** Either the program has a single global parameter or has multiple global parameters, we have a mapping from complexities, which are polynomials in terms of global parameters, to their scalar encodings. Since the final objective is the total complexity of the full transformed program, we need to sum the scalar encoding of complexities for all statements, without losing the ability to compare the resultants’ degrees. To that end, we propose to use a simple base-$|S|$ encoding method where $|S|$ is the maximum number of statements in the program: for a complexity encoded as scalar $c$, we use $|S|^c$ as a term in the final objective. As an example, to sum two complexities represented in scalar $c_1$ and $c_2$, we compute $|S|^{c_1} + |S|^{c_2}$. We define the base-$|S|$ sum of $c_i$ as $\sum |S|^{c_i}$.

**Indicator variable.** In the formulation, we require indicator variables to indicate if ST is disabled along a certain face – in which case no complexity reduction should be applied for the corresponding domain. We can use the big-M method, a well-known ILP modeling trick [Nemhauser and Wolsey 1988], to encode an indicator variable $y \in \{0, 1\}$ for the constraint $x = 0$ so that $y = 1$ iff $x = 0$.

5.3 Discussion

The above formulation is an integer objective bilinear constrained program. The objective is linear because it is an affine combination of the indicator variables. The problem is bilinear constrained because: in the original ILP formulation scheduling, the dependence matrix (defined in Section 3.3.2) is multiplied by a vector of unknowns to form a linear constraint; however by introducing the unknown reuse vectors $\vec{r}_i$, the dependence matrix contains entries that depends on $\vec{r}_i$, thereby making the constraints bilinear.

**Enabling transformations.** The enabling transformations presented in the original SR paper [Gautam and Rajopadhye 2006] can be incorporated into our formulation by the use of binary decision variables and a technique of encoding logical constraints as integer linear constraints.

6 MSSR HEURISTIC ALGORITHM

The problem formulation we present in Section 5.2 is a full characterization of the MSSR problem. In this work we consider this formulation only as a specification instead of a complete solution — solving an integer linear objective bilinear constrained program is NP-hard. As far as we know, none of well-known solvers can solve this problem out of the box, though it is possible to reformulate such problem into mixed integer linear programming (MILP) [Gupte et al. 2013]. However, the size of the formulation (i.e. total number of constraints and number of variables) in Section 5.2 is proportional to the number of statements, number of faces per statement and the maximal complexity of the program – either one of which could potentially lead to exponential blow up in the size of the formulation. Further, our formulation of dependency resolution is based on an
The key insights that guide our algorithm are that 1) choosing any valid reuse vector for a given ST results in the same final algorithmic complexity for the program and 2) for any valid reuse vector, the direction itself or its negation adheres to the program’s original affine schedule of the LHS of the reduction. We demonstrate these two insights with the following lemmas.

**Lemma 6.1.** For any application of ST, the complexity decrease is always the same regardless of the actual choice of reuse vector.

**Proof.** For any ST application, the reduction’s n-dimensional domain \( \mathcal{P} \) is reduced to the two half shells \( \mathcal{P} - \mathcal{P}' \) and \( \mathcal{P}' - \mathcal{P} \). The two half shells decompose into convex polyhedral sets corresponding to all \((n-1)\)-dimensional faces of \( \mathcal{P} \). Further, for each decomposed convex polyhedral set, the thickness of the set, which is defined as the spanned width of the set orthogonal to its corresponding face, is a constant dependent solely on the ST’s reuse vector and the face’s orientation. Therefore, the cardinality of each decomposed polyhedral set is just the cardinality of the face multiplied by some constant. It then follows that for any two STs with two non-zero reuse vectors, their resultant residual reductions’ complexities are always the same, and equal the sum of the cardinalities of all the faces of \( \mathcal{P} \) multiplied by some constant. \( \square \)

Before introducing the next lemma, we first introduce an extended definition of scheduling functions. Recall that the scheduling function of a reduction statement is an affine function from the reduction’s domain to the timestamp. We extend the context of a scheduling function from a reduction statement to the LHS of a reduction in a given program as follows. First the program is augmented by adding to the program a new redirect statement \( A[\overline{x}] = \lambda'[\overline{\bar{x}}] \) with the same domain as the domain of \( A \), where \( \lambda' \) is a fresh symbol which replaces the LHS array \( A \) of the program. Then the scheduling function of the LHS of the reduction is simply the scheduling function of the newly introduced redirect statement of the LHS in the schedule of the augmented program.

**Lemma 6.2.** Given the affine schedule for the augmented program, then for any ST application on a reduction whose operator has an inverse and for any valid reuse vector \( \bar{r} \), either \( \bar{r} \) or \(-\bar{r}\) agrees with the schedule of the original program and does not introduce a dependency cycle.

**Proof.** Consider a reduction statement \( S \) with projection \( \text{proj} \) and LHS array \( A \). Suppose \( A \) has an affine schedule \( \Theta_A \) then we have that \( \Theta_A \cdot [\overline{x}, \overline{\bar{p}}, 1]^T \) is the schedule time for \( A[\overline{x}] \). Let the vector \( \bar{r} \) be in the same space as the domain of \( S \), and we shift the domain of \( S \) along \( \bar{r} \); let the projected vector of \( \bar{r} \) onto the domain of \( \overline{A} \) be \( \overline{\bar{r}} = \text{proj}(\bar{r}) \). Consider \( \bar{x} \) and \( \overline{x} + \overline{\bar{r}} \). Their scheduled timestamps are \( \Theta_A \cdot [\overline{x}, \overline{\bar{p}}, 1]^T \) and \( \Theta_A \cdot [\overline{x} + \overline{\bar{r}}, \overline{\bar{p}}, 1]^T \). Since \( \Theta_A \cdot [\overline{x} + \overline{\bar{r}}, \overline{\bar{p}}, 1]^T - \Theta_A \cdot [\overline{x}, \overline{\bar{p}}, 1]^T = \Theta_A \cdot [\overline{\bar{r}}, 0, 0]^T \) is a constant not dependent on \( \overline{x} \), it must be the case that for all \( \overline{x} \), either \( A[\overline{x}] \) is always scheduled before \( A[\overline{x} + \overline{\bar{r}}] \), or vice versa. Specifically, if the first non-zero entry (in accordance with the timestamp comparison in Definition 3.5) of \( \Theta_A \cdot [\overline{\bar{r}}, 0, 0] \) is positive, then \( A[\overline{x}] \) is always scheduled
before $A[\vec{x} + \vec{r}]$; otherwise, $A[\vec{x}]$ is always scheduled after $A[\vec{x} + \vec{r}]$. If $A[\vec{x}]$ is scheduled before $A[\vec{x} + \vec{r}]$, then applying ST with reuse vector $\vec{r}$ will not introduce any dependence cycle, since the newly introduced dependency is always consistent with the original schedule; on the other hand, if $A[\vec{x}]$ is scheduled after $A[\vec{x} + \vec{r}]$, then applying ST with reuse vector $-\vec{r}$ will not introduce any dependence cycle.

Further, since $\vec{r}$ chosen this way is always consistent with the original schedule, a previous application of ST will not affect a later application of ST — intuitively, a previously applied ST introduces a dependency that can be subsumed by an enforced dependency according to the original program’s schedule; thus later a application of ST, as long as it is also consistent with original schedule, will not be affected.

\[\square\]

6.2 Algorithm

With justification in Section 6.1, we now introduce the heuristic algorithm in Figure 6.

(1) Schedule the augmented program to obtain an initial schedule $\Theta$ for all statements and LHS of reductions

(2) Apply ST to all faces of all reduction statement’s domains; choose the direction that is consistent with $\Theta$ by:

(a) First pick any valid reuse vector $\vec{r}$ from the candidate set.

(b) Test if $\vec{r}$ is consistent with $\Theta$, if not consistent, set $\vec{r} \leftarrow -\vec{r}$, if $-\vec{r}$ is also a valid reuse vector; otherwise, do not apply the current ST.

Fig. 6. SSSR heuristic algorithm

To test if $\vec{r}$ is consistent with $\Theta$, one can compute $\Theta_A \cdot \vec{r}_A$, with $\Theta_A$ and $\vec{r}_A$ defined as in Lemma 6.2, and then test if the first non-zero entry is positive. As an alternative way, one may attempt to reschedule the original augmented program with the introduced dependency along $\vec{r}$, and then test if the program is schedulable.

6.3 Algorithm Analysis

Heuristic scheduling. One advantage of the heuristic algorithm in Figure 6 is that the schedule $\Theta$ does not need to be obtained from forming and solving the ILP formulation as in Section 3.3, and one is free to choose any scheduling algorithm in the polyhedral literature such as [Bondhugula et al. 2008; Feautrier 1992a,b; Gupta et al. 2007]. Most of these algorithms, such as the PLUTO scheduler [Bondhugula et al. 2008] provide a scalable solution to the polyhedral scheduling problem and thus the algorithm in Figure 6 does not present bottleneck due to scheduling.

Optimality Guarantee. The algorithm is optimal for the MSSR problem if all the reduction operators have inverses. This is because the algorithm considers a basis direction of reuse, and picks the direction along that basis that is consistent with the original schedule. As long as all reduction operators have inverses, the heuristic algorithm will assign a non-zero reuse vector to each face that has valid reuse opportunities. In other words, the heuristic algorithm maximizes the total number of ST applications among all faces, if all reduce operators have inverses. For any application of ST along a face, the complexity decrease is always the same regardless of the actual choice of reuse vector. Therefore, maximizing the number of ST applications among all faces minimizes the total complexity.

Lastly, if a reduction operator does not have an inverse, thereby restricting the candidate set of directions, then it is possible for our algorithm to produce a non-optimal solution. Specifically, if
an operator does not have an inverse, the valid reuse vector for that operator will be restricted to a one sided direction (since ST requires the reuse direction to point out of certain boundaries of the polyhedral domain if the operator does not have an inverse), instead of both directions of the basis. It is possible that the original program does not have an unique valid schedule. Consider the following scenario: one schedule is consistent with \(\vec{r}\), while another schedule is consistent with \(-\vec{r}\); since the operator does not have an inverse, only the positive direction \(\vec{r}\) is valid. Therefore, the initial schedule will affect whether this ST is applied or not – which in turn leads to the suboptimality of the algorithm.

7 IMPLEMENTATION

We implemented our IR as in Section 3.2 and the heuristic algorithm as in Section 6 using Python. We used Integer Set Library (ISL) [Verdoolaege 2010] for manipulation of polyhedral set and relations. To obtain the original program schedule, we used a PLUTO-like scheduler built-in of ISL. To test if a reuse vector is consistent with the original schedule, we simply attempted to introduce a new dependency along the reuse vector and perform a full scheduling — note that this is not necessary, and can be potentially eliminated by the method of computing \(\Theta \cdot \vec{r}\) following Section 6.2. However, in our case simply attempting to reschedule the program is easier to implement and the method is agnostic to the underlying scheduling algorithm.

8 EVALUATION

The algorithm presented in this work is particularly effective on optimizing unoptimized implementations of probabilistic inference procedures into efficient implementations, where the inference procedures have mathematical specifications that naturally translate to our IR. The inference procedures are also iterative, so they contain multiple-statement reductions that are not addressed by previous work [Gautam and Rajopadhye 2006].

**Research Question.** The goal of this section is to evaluate how effective the heuristic algorithm is on improving the performance on benchmarks consisting of algorithms described above.

8.1 Evaluation Metrics

We considered the following two aspects to evaluate the effectiveness of the heuristic algorithm.

8.1.1 Complexity. We first evaluate asymptotic complexity of the algorithms. This is an important metric because it determines how well these algorithms scale to large data sets, and hence how widely they can be applied.

Optimality is defined regarding programs realizable through transformations presented in this work. We evaluated our implementation in Section 7 using unoptimized implementations of probabilistic inference procedures. We present their algorithmic complexities before and after optimization, as well as the optimal complexities achievable with transformations in this work, by solving the problem formulation in Section 5.2 exactly. In addition, we also report the complexities of manual implementations using transformations that are not in this work.

We collected the complexities before and after by counting the cardinality of the resultant polyhedral domains using library implementations in [Verdoolaege et al. 2007]. We collected the optimal complexities by inspecting the benchmarks and deriving the optimal complexities manually. We collected complexities of manual implementations by either finding an existing implementation of the algorithm if one exists in the literature or, otherwise, by manually deriving them.

8.1.2 Runtime. Note that a real performance gain is not necessarily implied by a complexity improvement because the asymptotic complexity comparison ignores constant factors. The constant factors can be caused by, for example, auxillary variables overhead or memory/cache effect induced
by ST, and the constant factors could change. Therefore, we further validate the runtime performance
gains due to lowered complexities, by measuring wall-clock time improvements between the
optimized and unoptimized implementations.

8.2 Benchmarks
A subset of the benchmark algorithms considered are identified as “model-algorithm” pairs, where
the model refers to a generative probabilistic model, and the algorithm refers to a class of algorithm
to perform inference on the model. We considered 3 models and 3 algorithms. For models, we con-
sidered the Gaussian Mixture Model (GMM) [Murphy 2012], Latent Dirichlet Allocation (LDA) [Blei
et al. 2003] and Dirichlet Multinomial Mixture (DMM) [Holmes et al. 2012]. For algorithms, we
considered Gibbs Sampling (GS) [Geman and Geman 1984], Metropolis Hasting (MH) [Hastings
1970; Metropolis et al. 1953] and Likelihood Weighting (LW) [Fung and Chang 1989]. Thus a total
of 9 “model-algorithm” pairs were considered as benchmark algorithms.

Models and algorithms above have broad applications in the literature. The models for LDA [Blei
et al. 2003; Griffiths and Steyvers 2004] and DMM [Holmes et al. 2012] are popular for existing data
science problems. The models for GMM [Daniel Huang 2017; Walia et al. 2018], LDA [Daniel Huang
2017; Walia et al. 2018], and DMM [Walia et al. 2018] have been used as benchmarks for probabilistic
inference systems. Gibbs sampling [Geman and Geman 1984], Metropolis-Hastings [Hastings 1970;
Metropolis et al. 1953], and Likelihood Weighting [Fung and Chang 1989] are all widely used
inference algorithms in the literature. LDA and DMM are particularly valuable benchmarks because
there are published Gibbs sampling algorithms that researchers have manually optimized ([Griffiths
and Steyvers 2004] and [Resnik and Hardisty 2010], respectively).

In addition, we included two other benchmarks, namely the Loopy Belief Propagation on a 2D
grid model for the application of Stereo matching (LBP-Stereo [Grauer-Gray and Cavazos 2011;
Jian Sun et al. 2003] ), and the Cox proportional hazards model (CoxPH) [Cox 1972; Therneau
2013]. Loopy Belief Propagation [Bishop 2006] is an iterative approximate inference algorithm, and
its instantiation on the 2D grid model has applications in fields such as vision [Grauer-Gray and
Cavazos 2011; Jian Sun et al. 2003] and physics [Kikuchi 1951]. CoxPH is a well known statistical
model, which is typically combined with Newton’s method, an iterative optimization algorithm, for
inference on the model’s parameters. CoxPH is commonly found in medical applications [Collett
1993; White et al. 2016], and mechanical systems [Susto et al. 2015].

All of the benchmarks have a common feature that they are iterative methods specialized to a
generative probabilistic model. The parameters for these benchmarks are listed in Table 1.

8.3 Results
As described in Section 8.1, we first evaluated our method by analyzing the asymptotic performance
improvements and reported the results in Section 8.3.1, then we validated the runtime improvements
of the benchmarks in Section 8.3.2.

8.3.1 Complexity Results. Table 2 summarizes the results on comparison of complexities, ex-
pressed in terms of the corresponding parameters of each benchmark.

The column “Original” gives the complexity of the original program for the benchmarks. The
column “Optimized (Heuristic)” (later abbreviated as “Optimized”) gives the complexity of the
transformed program using the heuristic implementation in Section 7. The column “MSSR-Optimal”
gives the complexity of the transformed program by potentially solving the problem formulation in
Section 5.2 exactly; and this is the optimal complexity one can achieve using techniques presented
in this work. The column “Manual” gives the complexity of a potential optimized manual imple-
mentation written by a developer; this means that the complexity reduction potentially comes from
| Benchmark(s) | Parameter | Meaning                  |
|-------------|-----------|--------------------------|
| GMM-*       | N         | number of observations   |
|             | K         | number of clusters       |
| LDA-*       | W         | number of words in corpus|
|             | K         | number of topics         |
|             | W         | number of words in corpus|
|             | K         | number of topics         |
| DMM-*       | D         | number of documents in corpus|
|             | A         | size of alphabet of corpus|
|             | L         | maximum length of document|
|             | N         | number of total pixels    |
|             | K         | maximum stereo displacement|
|             | D         | number of neighbors of a pixel|
|             | N         | number of observations    |
|             | K         | dimension of a single observation|

Table 1. Benchmarks: Parameters and Meanings

| Benchmark | Original | Optimized (Heuristic) | MSSR-Optimal | Manual | #IR |
|-----------|----------|-----------------------|--------------|--------|-----|
| GMM-GS    | $O(N^2K^2)$ | $O(NK)$               | $O(NK)$      | $O(NK)$ | 16  |
| GMM-MH    | $O(N(N + K))$ | $O(N)$                | $O(N)$       | $O(N)$ | 16  |
| GMM-LW    | $O(N(N + K))$ | $O(N)$                | $O(N)$       | $O(N)$ | 5   |
| LDA-GS    | $O(W^2K^2)$ | $O(WK)$               | $O(WK)$      | $O(WK)$ | 20  |
| LDA-MH    | $O(W^2K)$   | $O(WK)$               | $O(WK)$      | $O(WK)$ | 42  |
| LDA-LW    | $O(W^2K)$   | $O(W)$                | $O(W)$       | $O(W)$ | 7   |
| DMM-GS    | $O(WADK^2 + D^2K^2)$ | $O((W + A)KD)$       | $O((W + A)KD)$ | $O(AKD)$ | 40  |
| DMM-MH    | $O(D^2K^2 + D(W + A))$ | $O((K + W + A)D)$ | $O((K + W + A)D)$ | $O((K + L + A)D)$ | 82  |
| DMM-LW    | $O((W + K)D)$ | $O((K + W + A)D)$ | $O((K + W + A)D)$ | $O((K + L + A)D)$ | 10  |
| LBP-Stereo| $O(NK^2D^2)$ | $O(NKD)$              | $O(NKD)$     | $O(NKD)$ | 3   |
| CoxPH     | $O(K^2N^2)$ | $O(K^2N)$             | $O(K^2N)$    | $O(K^2N)$ | 6   |

Table 2. Benchmarks: Comparison of Complexities

transformations not present in this work. The column “#IR” counts the number of IR statements for the benchmark.

Comparing the “Original” and “Optimized” columns, our approach can reduce the complexity for all benchmarks. Comparing the “Optimized” and “MSSR-Optimal” columns, our approach can generate algorithms that have the same complexity as that of optimal implementation for all benchmarks. Comparing the “Optimized” and “Manual” columns, our approach can generate algorithms with complexities the same as manual implementations for 8 out of 11 benchmarks. We identified that the 3 benchmarks related to DMM require additional data layout modifications which we did not consider in this work, which is a direction of future research.

8.3.2 Runtime Validation. So far we have evaluated our heuristic algorithm using algorithmic complexity as the primary factor, which ignores constant factors caused by, for example, auxiliary variables overhead or memory/cache effect induced by ST. In this section, we validate our hypothesis that asymptotic complexity improvements dominates potential constant factors improvements for the parameters of these benchmarks by timing our benchmarks and comparing the runtimes of the naive implementations with the optimized implementations. We use C implementations that
match the naive and optimized complexities reported in Table 2. We ran these implementations and report timeouts for benchmarks that ran for 12+ hours.

**Parameter sizes.** We collected the typical instantiated values for global parameters from the corresponding literatures. Specifically, for GMM we use [Daniel Huang 2017], for LDA we use [Newman 2008], for DMM we use [Turnbaugh et al. 2008], for LBP-Stereo we use [Jian Sun et al. 2003] and for CoxPH we use [Therneau 2013, Appendix D2]. Based on these prior works, we collected the following parameters for each model in Table 3.

**Results.** In Table 4, all benchmarks have non-trivial speedups. In particular, for LDA benchmarks all the unoptimized implementation timeout. This is because, in Table 2, complexities of LDA benchmarks all improve by a factor of $W \times K$. With our instantiated parameter values this factor is $466,000 \times 50 = 2.3 \times 10^7$ – the largest factor across all benchmarks; this large a factor unsurprisingly leads to timeout of the unoptimized implementations. On the other hand, in Table 2 LBP-Stereo’s complexity only improves by a factor of $D$, the number of neighbors of a pixel, which is set to 4 (i.e. number immediate neighbors of a pixel) in our parameter setting. Nonetheless, we observe a speedup of 1.1x for this benchmark. We also note that this speedup scales with the specification of the LBP model – setting to 8 neighboring pixels (i.e. nearby 8 pixels for a pixel at the center of a 3 by 3 square) would lead to speed up of of 1.4x. In summary, the observed speedups validates that for these benchmarks and our technique, complexity dominates constant-factor concerns.

| Benchmark(s) | Parameter | Size |
| --- | --- | --- |
| GMM-* | $N$ | 10000 |
| | $K$ | 10 |
| LDA-* | $W$ | 466,000 |
| | $K$ | 50 |
| DMM-* | $W$ | 570,000 |
| | $K$ | 4 |
| | $D$ | 278 |
| | $A$ | 129 |
| | $L$ | 3202 |
| LBP-Stereo | $N$ | 110,592 |
| | $K$ | 16 |
| | $D$ | 4 |
| CoxPH | $N$ | 424 |
| | $K$ | 11 |

Table 3. Benchmarks: Parameter Sizes

| Benchmark | Original | Optimized | Speedup |
| --- | --- | --- | --- |
| GMM-GS | 20.8 ms | 2.8 ms | 7.4 x |
| GMM-MH | 0.49 ms | 0.37 ms | 1.3 x |
| GMM-LW | 2.4 s | 1.6 s | 1.5 x |
| LDA-GS | **timeout** | 3.1 ms | > $14 \times 10^6$ x |
| LDA-MH | **timeout** | 0.53 s | > $82 \times 10^3$ x |
| LDA-LW | **timeout** | 36.2 s | > $1.2 \times 10^3$ x |
| DMM-GS | 2.2 s | 0.54 s | 4.1 x |
| DMM-MH | 389 ms | 9 ms | 43 x |
| DMM-LW | 48 s | 0.78 s | 62 x |
| LBP-Stereo | 14.9 s | 13.8 s | 1.1 x |
| CoxPH | 54.1 ms | 6.5 ms | 8.3 x |

Table 4. Runtime evaluations

9 RELATED WORK

**Simplifying Reductions.** Previous works on simplifying reductions are [Liu et al. 2005] and [Gautam and Rajopadhye 2006]. Liu et al. [2005] proposed a loop based transformation algorithm for reducing complexities on loop programs. The algorithm uses the Omega calculator [Padua 2011] for analysis on a contributing set. The method is general in that any set calculation method, potentially methods that work for even non-polyhedral sets, can be used. The method in Liu et al. [2005] uses only the direction of loop increment to decrease the complexity. Gautam and Rajopadhye [2006] generalized the method in [Liu et al. 2005]; one of the advances was that it formalized the notion of reuse space and proposed to use directions in the reuse space to decrease complexity.
Incrementalization in Probabilistic Programming. The problem of incrementalization occurs in probabilistic programming system (PPS), and is known as incremental inference. Existing work such as [Kiselyov 2016; Nori et al. 2015; Ritchie et al. 2016; Wu et al. 2016; Yang et al. 2014; Zhang and Xue 2019] attempt to address the problem of incremental inference in PPS. However, these techniques are variants/combinations of 1) tracing JITs, 2) specialization and caching/mem- oization, 3) dynamic dependence analysis, 4) dynamic program slicing, or 5) runtime symbolic analysis – in summary, dynamic optimizations. These techniques introduce significant runtime overhead for storing dependency graph/traces (which is of size proportional to the number of the executed statement instances) and/or performing analysis on those graphs/traces dynamically. Our technique can be applied to PPS to solve the incremental inference problem; however, our technique is a static compilation technique which do not suffer from runtime overhead.

Many existing and ongoing work [Atkinson et al. 2018; Bingham et al. 2018; Daniel Huang 2017; Goodman et al. 2008; Goodman and Stuhlmüller 2014; Mansingkha et al. 2018; Narayanan et al. 2016; Plummer 2015] allow the user to code in high level DSLs. Though the details on these systems are out of the scope of the paper, our method can be potentially integrated into these systems for generating code with efficient complexity.

Reductions. Previous work [Doerfert et al. 2015; Ginsbach and O’Boyle 2017] proposed techniques to detect reductions from loop based code; these techniques can be used as front-ends to our technique for conversion into our reduction based IR. Previous work [Doerfert et al. 2015; Ginsbach and O’Boyle 2017; Rauchwerger and Padua 1999; Reddy et al. 2016] optimizes reductions in the polyhedral model for considerations such as privatization and parallelization. They do not optimize reductions’ complexities; however, they can be used as optimizing backends for generating efficient code for reductions after applying our method.

ILP scheduling. Previous work [Pouchet et al. 2008, 2007, 2011] gives an ILP formulation of the scheduling problem. Specifically, [Pouchet et al. 2011] shows how to construct constraints for a convex ILP problem to find an $m$-dimensional schedule for a program. Moreover, this formulation of constraints allows one to incorporate a desired objective to be optimized — in this work, we used the complexity of the final transformed program as the objective and we showed how to encode such objective as a affine expression in Section 5.2.3.

Heuristic scheduling. There are also other scheduling methods such as the ones in [Bondhugula et al. 2008; Feautrier 1992a,b] that use heuristics to schedule a program. These methods are usually more scalable than an ILP formulation. In this work, we use ideas from the ILP formulation to formulate the MSSR problem, while our provided heuristic algorithm does not depend on using the ILP formulation for scheduling. Instead, we use the PLUTO [Bondhugula et al. 2008; Verdoolaege 2010] heuristic scheduling algorithm in our implementation.

10 CONCLUSION

In this work, we introduce the multiple-statement reduction problem and provide a heuristic algorithm that is optimal for reduction operators that have inverses. These reductions have otherwise only appeared as domain- or algorithm-specific optimizations as described in the published description of standard probabilistic inference algorithms. Our hope is that this work formally outlines a key general-purpose optimization opportunity that can be delegated to the compiler, rather than being a significant piece of manual implementation that stands between the elaboration of a new probabilistic inference algorithm and its high performance implementation. Our results hold the promise that emerging languages and systems for this increasingly important class of computations could see significant performance improvements by incorporating our techniques.
REFERENCES

Eric Atkinson, Cambridge Yang, and Michael Carbin. 2018. Verifying Handcoded Probabilistic Inference Procedures. In arXiv e-prints.

Mohamed-Walid Benabderrahmane, Louis-Noël Pouchet, Albert Cohen, and Cédric Bastoul. 2010. The Polyhedral Model is More Widely Applicable Than You Think. In European Conference on Theory and Practice of Software, International Conference on Compiler Construction.

Eli Bingham, Jonathan P Chen, Martin Jankowiak, Fritz Obermeyer, Neeraj Pradhan, Theofanis Karaletsos, Rohit Singh, Paul Szerlip, Paul Horsfall, and Noah D Goodman. 2018. Pyro: Deep Universal Probabilistic Programming. arXiv preprint arXiv:1810.09538 (2018).

Christopher M. Bishop. 2006. Pattern Recognition and Machine Learning (Information Science and Statistics). Springer-Verlag, Berlin, Heidelberg.

David M. Blei, Andrew Y. Ng, and Michael I. Jordan. 2003. Latent Dirichlet Allocation. In JMLR, Vol. 3.

Uday Bondhugula, Albert Hartono, J. Ramanujam, and P. Sadayappan. 2008. A Practical Automatic Polyhedral Parallelizer and Locality Optimizer. In Programming Language Design and Implementation.

Jean-François Collard, Denis Barthou, and Paul Feautrier. 1995. Fuzzy Array Dataflow Analysis. In ACM SIGPLAN Symposium on Principles and Practice of Parallel Programming.

D Collett. 1993. Modelling Survival Data in Medical Research. New York: Chapman and Hall/CRC.

D. R. Cox. 1972. Regression Models and Life Tables. Journal of the Royal Statistical Society: Series B (Methodological) 34, 2 (1972), 187–202.

Marco F Cusumano-Towner, Feras A Saad, Alexander K Lew, and Vikash K Mansinghka. 2019. Gen: a general-purpose probabilistic programming system with programmable inference. In Programming Language Design and Implementation. Greg Morisset Daniel Huang, Jean-Baptiste Tristan. 2017. Compiling Markov Chain Monte Carlo Algorithms for Probabilistic Modeling. In Programming Language Design and Implementation.

Johannes Doerfert, Kevin Streit, Sebastian Hack, and Zino Benaisa. 2015. Polly’s Polyhedral Scheduling in the Presence of Reductions. In International Workshop on Polyhedral Compilation Techniques.

E. Ehrhardt. 2009. Sur un probblème de géométrie diophantienne linéaire. II. Journal für die reine und angewandte Mathematik 1967 (12 2009). https://doi.org/10.1515/crll.1967.227.25

P. Feautrier. 1988. Array Expansion. In International Conference on Supercomputing.

Paul Feautrier. 1992a. Some efficient solutions to the affine scheduling problem. I. One-dimensional time. International Journal of Parallel Programming 21, 5 (01 Oct. 1992), 313–347. https://doi.org/10.1007/BF01407835

Paul Feautrier. 1992b. Some efficient solutions to the affine scheduling problem. Part II. Multidimensional time. International Journal of Parallel Programming 21, 6 (01 Dec. 1992), 389–420. https://doi.org/10.1007/BF01379404

Robert M. Fung and Kuo-Chu Chang. 1989. Weighing and Integrating Evidence for Stochastic Simulation on Bayesian Networks. In UAI.

Gautam and S. Rajopadhye. 2006. Simplifying Reductions. In ACM SIGPLAN Symposium on Principles of Programming Languages.

Andrew Gelman, Daniel Lee, and Jiqiang Guo. 2015. Stan: A probabilistic programming language for Bayesian inference and optimization. Journal of Educational and Behavioral Statistics 40, 5 (2015), 530–543.

Stuart Geman and Donald Geman. 1984. Stochastic Relaxation, Gibbs Distributions, and the Bayesian Restoration of images. In IEEE Transactions on Pattern Analysis and Machine Intelligence.

Philip Ginsbach and Michael F. P. O’Boyle. 2017. Discovery and Exploitation of General Reductions: A Constraint Based Approach. In International Symposium on Code Generation and Optimization.

Noah D. Goodman, Vikash K. Mansinghka, Daniel M. Roy, Keith Bonawitz, and Joshua B. Tenenbaum. 2008. Church: A language for generative models. In UAI.

Noah D Goodman and Andreas Stuhlmüller. 2014. The Design and Implementation of Probabilistic Programming Languages. Accessed: 2016-10-7.

Scott Grauer-Gray and John Cavazos. 2011. Optimizing and Auto-tuning Belief Propagation on the GPU. In Languages and Compilers for Parallel Computing. Springer Berlin Heidelberg.

T. Griffiths and M. Steyvers. 2004. Finding Scientific Topics. In PNAS, Vol. 101.

Gautam Gupta, Kim Daegon, and Sanjay Rajopadhye. 2007. Scheduling in the Z-Polyhedral Model. International Parallel and Distributed Processing Symposium.

Akshay Gupta, Shabbir Ahmed, Myun Cheon, and Santanu Dey. 2013. Solving Mixed Integer Bilinear Problems Using MILP Formulations. SIAM Journal on Optimization 23 (04 2013). https://doi.org/10.1137/110836183

W. K. Hastings. 1970. Monte Carlo Sampling Methods Using Markov Chains and Their Applications. In Biometrika, Vol. 57. Ian Holmes, Keith Harris, and Christopher Quince. 2012. Dirichlet Multinomial Mixtures: Generative Models for Microbial Metagenomics. In PLOS One.
Rajan Walia, Jacques Carette, Praveen Narayanan, Chung-chieh Shan, and Sam Tobin-Hochstadt. 2018. Efficient Compilation of Array Probabilistic Programs. In arXiv e-prints.

Nicola White, Fiona Reid, Adam Harris, Priscilla Harries, and Patrick Stone. 2016. A Systematic Review of Predictions of Survival in Palliative Care: How Accurate Are Clinicians and Who Are the Experts? PLOS ONE 11, 8 (08 2016), 1–20. https://doi.org/10.1371/journal.pone.0161407

Yi Wu, Lei Li, Stuart Russell, and Rastislav Bodík. 2016. In International Joint Conferences on Artificial Intelligence.

Lingfeng Yang, Patrick Hanrahan, and Noah Goodman. 2014. Generating Efficient MCMC Kernels from Probabilistic Programs. In Proceedings of the Seventeenth International Conference on Artificial Intelligence and Statistics.

Tomofumi Yuki, Gautam Gupta, DaeGon Kim, Tanveer Pathan, and Sanjay Rajopadhye. 2013. AlphaZ: A System for Design Space Exploration in the Polyhedral Model. In Languages and Compilers for Parallel Computing, Hironori Kasahara and Keiji Kimura (Eds.). Springer Berlin Heidelberg, Berlin, Heidelberg, 17–31.

Jieyuan Zhang and Jingling Xue. 2019. Incremental Precision-Preserving Symbolic Inference for Probabilistic Programs.
**A EXTRA LISTINGS**

```plaintext
1 for( i = 0; i < N; i++)
2 B[N - 1] += A[i]
3 for( i = N-2; i >= 0; i--)
4 B[i] = B[i+1] - A[i]
```

Listing 7. Alternative optimized PS (right-to-left)

---

**B SIMPLIFYING REDUCTION**

A key opportunity that we’ve identified is the integration of the histogram transformation with the Simplifying Reduction transformation [Gautam and Rajopadhye 2006].

Consider an IR statement for which the set of non-affine equality predicates Q is empty:

1 label: LHS[u] ⊕= expr : ∅ & P

Simplifying reduction (SR) transforms this statement into an equivalent form as in Figure 7. The transformation takes in one parameter, a nonzero constant vector \( \vec{r} \), representing the direction of reuse, which we will explain shortly.

We first define some notations:

- we use \( \rho^a \) to denote projecting \( p \) onto space \( a \); the superscript acts effectively as a projection function; \( p \) can either be a point, an affine transformation or a polyhedral set of points.
- \( T_r(x) \) is an affine translation transformation (under homogeneous coordinates). That is, if \( x \) is a vector \( \vec{x} \) representing a point, \( T_r \) shifts \( \vec{x} \) to \( \vec{x} + \vec{r} \). If \( x \) is a polyhedron \( P \), \( T_r \) shifts all points in \( P \) by \( +\vec{r} \).

Then, let \( P' = T_{-\vec{r}}(P) \), i.e. \( P' \) is \( P \) shifted by \( -\vec{r} \), we define the following symbols in Figure 7:

\[
\begin{align*}
P_{\text{add}} &= P - P' \quad P_{\text{sub}} = P' - P \quad P_{\text{int}} = P \cap P' 
\end{align*}
\]

**Explanation.** The core intuition behind ST is to realize *reuse* of the RHS \( expr \). Specifically, we require a choice of \( \vec{r} \) so that it presents sharing for the RHS expression, that is:

\[
\llbracket \text{sub}(\text{expr}, T_r(\text{freevars(expr)})) \rrbracket = \text{expr}
\]

In other words, the value of \( expr \) is the same for any point \( v \) and its shifted counterpart \( T_r(v) \). This way, we can avoid evaluation of \( expr \) by simply copying from \( \text{sub}(\text{expr}, T_r(\text{freevars(expr)})) \), whenever possible. The first five statements 1-add-only through 1-add-reuse-sub computes LHS this way and reuse \( \text{sub}(\text{expr}, T_r(\text{freevars(expr)})) \) along \( \vec{r} \). The domains of the five statements prescribe the set of points according to each statement’s semantics.
- \(P_{\text{add}}^u\) is the set of indices that receive \(\text{expr}\)'s values evaluated in \(P_{\text{add}}\).
- \(P_{\text{sub}}^u\) is the set of indices that receive \(\text{expr}\)'s values evaluated in \(P_{\text{sub}}\).
- \(P_{\text{int}}^u\) is the set of indices that receive \(\text{expr}\)'s values evaluated in \(P_{\text{int}}\).

Receiving value from the intersection means that it is possible to reuse from the index point shifted by \(\bar{r}\).

We then explain each of the first five statements in turn:

**Reuse only.** Consider the domain of 1-reuse-only, \(P_{\text{int}}^u - (P_{\text{add}}^u \cup P_{\text{sub}}^u)\), can be read as: the set of indices that receive value from intersection, but does not receive from ADD or SUB, and this is precisely the set of points that can be directly copied along \(\bar{r}\). Thus, 1-reuse-only performs just this copy operation: \(LHS[u] = LHS[T^u_r(u)]\).

**Add Only.** 1-add-only's domain \(P_{\text{add}}^u - P_{\text{int}}^u\) can be read as: the set of indices that receive value from ADD, but does not receive value from intersection. One can verify that \(P_{\text{add}}^u - P_{\text{int}}^u = P_{\text{add}}^u - P_{\text{int}}^u - P_{\text{sub}}^u\), this also implies that the set also does not receive value from SUB. Therefore, the statement just copies from ADD.

**Add and Reuse.** 1-add-reuse's domain, \(P_{\text{add}}^u \cap (P_{\text{int}}^u - P_{\text{sub}}^u)\), can be read as: the set of indices that receive value from ADD and the intersection, but does not receive value from SUB. Therefore the statement reuses value along \(\bar{r}\), and increments with value calculated from ADD.

**Sub and Reuse.** 1-reuse-sub's domain, \(P_{\text{sub}}^u \cap (P_{\text{int}}^u - P_{\text{add}}^u)\), can be read as: the set of indices that receives value from SUB and the intersection, but does not receive value from ADD. Therefore the statement reuses value along \(\bar{r}\), and decrements with value calculated from SUB.

**Add, Reuse and Sub.** 1-add-reuse-sub's domain, \(P_{\text{sub}}^u \cap P_{\text{int}}^u \cap P_{\text{add}}^u\), can be read as: the set of indices that receive value from both ADD, the intersection and SUB. Therefore, the statement reuse along \(\bar{r}\), increments with ADD and decrements with SUB.

**Residual Reductions.** The statements 1add and 1sub are themselves reductions, and we will call them residual reductions after SR transformation. They compute additional values that are requested by the top five statements. The residual reduction accumulates the same right-side expression as the original reduction, but with domains that are subsets of the original domain.

### B.2 Configuration of Simplifying Reduction

As mentioned in Section 4.2, we need to consider three constraints – complexity, sharing and dependence – when choosing the reuse vector \(\bar{r}\). Here we discuss the constraints in more detail.

**Complexity.** A program's complexity is a function over its input parameters.

The complexity after one SR transformation is equal to the total sum of all cardinalities of domain sizes of the statements after the transformation. The complexity of the first five statements combined together is equivalent to iterating points of LHS array, and therefore it will always remain unchanged, since we will always need to compute answer for each point of the LHS. As shown in [Gautam and Rajopadhye 2006], in order for one step of SR to be meaningful, in the sense that it decreases the complexity, we need that \(|P_{\text{add}}| + |(P_{\text{int}}^u)' \cap P_{\text{sub}}| < |P|\).

**Sharing.** Fully determining all possible \(\bar{r}\) that presents sharing for the right-hand side expression is not decidable: for an arbitrary RHS expression \(\text{expr}\) as an uninterpreted function: we can encode the problem as \(\exists \bar{r} \forall v. \llbracket \text{expr} \rrbracket = \llbracket \text{subst(\text{expr}, T_r(freevars(\text{expr})))} \rrbracket\), and this is not decidable in general. However, we can still heuristically deduce valid reuse vectors, if we know the internals structure of \(\text{expr}\). [Gautam and Rajopadhye 2006] proposed a heuristical approach by computing a polyhedral
set $\hat{S}_2$ called share space, which is the intersection of the nullspaces of the dependence functions of all the subexpressions of $expr$, and selecting any vector $r \in \hat{S}_2$.

B.3 Choosing a reuse vector

Complexity reduction. We chose $r$ to be in the lineality space $L_P$, that is defined informally as the subspace of $P$ where $P$ extends infinitely as the sizes the parameters of $P$ tends to infinity. Intuitively, this means that we only want to reuse computation along directions of $P$ that can grow asymptotically with the parameters, instead of directions that are bounded by fixed constants.

Sharing. We use a simple yet effective heuristic in our implementation: find the set of variables $fv$ in the left hand side $u$ that is not bound in the right hand side $v$: i.e. $fv = s - v$ (recall that $s$ is the space of the statement’s domain $P$). For any variable $x \in fv$, we can find its constant unit vector $\hat{r}_x$ (a vector under space $s$ that is 1 only along direction $x$), we must have $\hat{r}_x$ satisfying the above criterion: $\forall v. expr = expr(T_{\hat{r}_x}(v))$. In fact, any linear scaling of $\hat{r}_x$ is a valid choice. In summary, for one IR statement, there can be $|fv|$ dimensions of reuse, in the sense that each $\hat{r}_x \forall x \in fv$ are orthogonal to each other and thus forms different dimensions; for each dimension, there can be infinitely many valid reuse vectors, that are the different scalings of the unit direction for that dimension.

Dependence. Applying SR can introduce new dependencies along the reuse vector that was not in the original program. For example, in Listing 6, the transformed statement T-add-reuse introduces a new dependence that now $S[i]$ depends on $S[i - 1]$. We require that applying SR along a vector $r$ does not introduce any dependency cycle in transformed program so that it remains valid.

C ENABLING TRANSFORMATIONS

In this section we briefly review the enabling transformations introduced in [Gautam and Rajopadhye 2006]. Since these transformations are important to fully utilize ST for a single reduction, we encourage readers to find more details of these transformations in [Gautam and Rajopadhye 2006].

Reduction Decomposition. For reduction with projection function $proj$, we can potentially decompose $proj = proj1 \circ proj2$, where $\circ$ denotes function composition. It is possible to break the reduction into two statements: the first statement with projection $proj2$ produces an intermediate output, followed by a reduction with projection $proj1$ that returns the original output. The first statement could lead to a larger share space than the original reduction, and therefore RD enables enable ST.

Same Operator Transformation. It’s possible to lift inner expressions out of reductions to increase share space.

Distributivity Transformation. It’s possible to utilize distributivity of an operator to lift inner expression out of reductions to increase share space.

Higher Order Operator Transformation. It’s possible to collapse along the entire reuse space, if the reduce operator $\oplus$ has an higher order operator $\otimes$. 
