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Schenck, Robert; Rønning, Ola; Henriksen, Troels; Oancea, Cosmin E.

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AD for an Array Language with Nested Parallelism

Robert Schenck, Ola Rønning, Troels Henriksen and Cosmin E. Oancea

DIKU, University of Copenhagen, Denmark

r@bert.lol, ola@di.ku.dk, athas@sigkill.dk, cosmin.oancea@diku.dk

Abstract

We present a technique for applying (forward and) reverse-mode automatic differentiation (AD) on a non-recursive second-order functional array language that supports nested parallelism and is primarily aimed at efficient GPU execution.

The key idea is to eliminate the need for a “tape” by relying on redundant execution to bring into each new scope all program variables that may be needed by the differentiated code. Efficient execution is enabled by the observation that perfectly-nested scopes do not introduce re-execution, and such perfect nests are produced by known compiler transformations, e.g., flattening. Our technique differentiates loops and bulk-parallel operators—such as map, reduce, histogram, scan, scatter—by specific rewrite rules, and aggressively optimizes the resulting nested-parallel code. We report an experimental evaluation that compares with established AD solutions and demonstrates competitive performance on nine common benchmarks from recent applied AD literature.

Keywords: automatic differentiation, functional data parallel language, compilers, GPGPU.

1 Introduction

Automatic differentiation (AD) is a practical way for computing derivatives of functions that are expressed as programs. AD of sequential code is implemented in tools such as Tapenade [2], ADOL-C [14], and Stalingrad [35]. Modern deep learning is built on array programming frameworks such as Tensorflow [1] and PyTorch [32], which provide implicitly parallel bulk operations that support AD.

A largely-unsolved challenge is supporting AD for high-level parallel languages [18, 33, 44] that permit arbitrary nesting of sequential and parallel constructs. Such solutions may in principle act as a catalyst for prototyping and training of more advanced machine learning (ML) models.

ML involves minimising a cost function, which typically has far more inputs than outputs. The reverse mode of AD is the most efficient in such cases [3], but is challenging to implement because intermediate program values are required by the differentiated code. The program must first run a forward sweep (a.k.a., primal trace) that stores intermediate program states on the tape, which is then read during the return sweep, which essentially runs the program in reverse, and actually computes the derivative.

A significant amount of work has studied how to elegantly model reverse mode AD as a compiler transformation, and how to hide the tape under powerful programming abstractions such as (dynamic) closures [35] and delimited continuations [46]. These abstractions are not suited for efficient parallel execution on manycore hardware such as GPUs.

This work is to our knowledge the first to demonstrate an efficient GPU implementation of reverse mode AD as a compiler transformation on a data-parallel language that supports nested parallelism by means of higher-order array combinators (SOAC), such as map, reduce(-by-index), scan.

A key difference is that related approaches save by default all variables on the tape and support checkpointing annotations as an optimization (of memory footprint). However, in a nested-parallel context, the tape may give raise to complex, irregular data-structures that are passed across deep nests and that are challenging to implement efficiently in regards to optimizing spatial (coalescing) and temporal locality.

In contrast, our modeling of the tape takes inspiration from the idea [3] that applying reverse AD to a straight line of side-effect-free code does not require any tape per se, because all intermediate values remain available. We expand this idea to drive the code-transformation across lexical scopes by requiring that whenever the return sweep enters a new scope s, it first redundantly re-executes the forward sweep of s in order to bring all the needed variables into scope.

Our technique preserves the work and span asymptotics because the recomputation overhead is at worst proportional to the depth of the deepest nest of scopes, which is constant for a given nonrecursive program. Moreover, perfectly-nested scopes (other than loops) are guaranteed to not introduce re-execution, hence the overhead can be minimized by classic compiler transformations such as flattening nested parallelism [19] and polyhedral-like optimizations [7]. Since our tape is essentially formed by the in-scope variables, scalars are efficiently accessed from registers rather than global memory (tape), and the code resulted from AD fully benefits from the existent compiler-optimization repertoire.

In what sequential loops are concerned, the loop-variant variables require checkpointing because the return sweep needs to execute the iterations in reverse order. In addition, we exploit to some extent the fundamental time-space trade-off studied by Siskind and Pearlmutter [42], in a simple and practical way, by allowing the user to annotate how many times a loop should be strip-mined.\(^1\)

\(^1\) Strip-mining \(k\) times a loop of count \(n\) requires only \(2k\times n\) more memory than the original program at the expense of \(k\times n\) re-execution at worst. Of course, since the strip-mining factor is a user-defined constant it cannot achieve the logarithmic time-space overhead.
Having designed the glue that binds scopes together, we turn our attention to deriving high-level rewrite rules for differentiating the parallel operators of the language. We achieve this by starting from the main rewrite rule of the reverse mode and extend it by applying reasoning that combines imperative (dependence analysis, loop distribution) and functional thinking (rewrite rules, recurrences as scans).

In particular, the simplest parallel operator, namely map, is the most difficult one to translate, because its purely functional semantics allows free variables to be freely read inside it, but reverse AD replaces a read with an accumulation, which cannot be in general represented as a combination of classical data-parallel constructs. In this context we report optimizations related to turning accumulators to more-specialized constructs, such as reductions and generalized histograms [17], that can yield speedups close to one order of magnitude at the application level (e.g., GMM and LSTM).

Our overall contribution is an end-to-end algorithm of AD with support for nested parallel combinators, implemented inside (a clone of) the compiler for the Futhark language and aimed at GPU execution. Our specific contributions are:

- A redundant-execution technique for reverse AD that (i) eliminates the need for tape, (ii) is guaranteed to not introduce re-execution for perfectly-nested scopes other than loops, and (iii) practically supports the time-space tradeoff by means of loop stripmining.
- A set of rewrite rules for differentiating higher-order parallel combinators, including uses of free variables.
- A collection of optimisations that rewrite common cases of accumulators to other specific language constructs that benefit from specialized code generation.
- An experimental evaluation that demonstrates (i) sequential performance competitive with Tapenade [2] on ADBench [43], (ii) GPU performance competitive with Enzyme [29] on two of their applications, i.e., RS-Bench, XS-Bench, and (iii) significant GPU speedups in comparison with PyTorch [32] on GMM, LSTM, and for the sparse formulation of K-Means.

2 Preliminaries

This section provides the gist of how automatic differentiation of programs may be derived from its mathematical foundation. Given a function \( P : \mathbb{R}^a \rightarrow \mathbb{R}^d \), the (total) derivative of \( P \) at a point \( x \in \mathbb{R}^a \) (where \( P \) is differentiable) can be written as a function of \( y \in \mathbb{R}^a \) in terms of its Jacobian as \( \frac{\partial P}{\partial x}(y) = J_P(x) \cdot y \), where \( \cdot \) is matrix-vector multiplication.

The Jacobian is a representation of a function derivative at a specific point. The chain rule provides a straightforward way of deriving the derivative across function composition.

\[ \frac{\partial f}{\partial a} \]

An original-program statement \( x = f(a) \) produces the differentiated code \( \nabla x = \frac{\partial f(a)}{\partial a} \cdot \nabla \), where \( \nabla \) denotes the adjoint of program variable \( x \).

Figure 1. (a) A program \( P \) which computes the function \( f(x_0, x_1) = (x_1 \cdot \sin(x_0), x_0 \cdot x_2) \), (b) the forward mode AD transformation of the program, and (c) the reverse mode AD transformation of the program.

For example, if \( P(x) = h(g(f(x))) \), and the types of \( f, g, h \) are \( f : \mathbb{R}^a \rightarrow \mathbb{R}^b, g : \mathbb{R}^b \rightarrow \mathbb{R}^c \), and \( h : \mathbb{R}^c \rightarrow \mathbb{R}^d \), then:

\[
J_{\dot{P}}(x) = J_h(g(f(x))) \cdot J_g(f(x)) \cdot J_f(x) \tag{1}
\]

Since matrix multiplication is associative, there are multiple ways to evaluate \( J_{\dot{P}} \) when it’s viewed as a program. For example, from right-to-left:

\[
\dot{J}_{\dot{P}}(x) = J_h(g(f(x))) \left[ J_g(f(x)) \right] J_f(x) \tag{2}
\]

or left-to-right:

\[
\dot{J}_{\dot{P}}(x) = J_h(g(f(x))) J_g(f(x)) J_f(x) \tag{2}
\]

Notice that if \( h \) is a scalar function (i.e., \( d = 1 \)), then \( J_h(g(f(x))) \) is a gradient vector of dimension \( \mathbb{R}^d \). In the right-to-left formulation, \( \star \) is a large matrix of dimension \( \mathbb{R}^{C \times A} \), whereas in the left-to-right formulation, \( \star \) is a vector of dimension \( \mathbb{R}^b \). When \( A \ll d \), it is much more expensive to compute/manifest the larger intermediate Jacobian matrices in the right-to-left evaluation order. When \( A \ll d \), it’s more efficient to choose the right-to-left evaluation order. There’s a catch: the Jacobians depend on intermediate results, e.g., \( J_g(f(x)) \) depends on \( f(x) \). In the right-to-left evaluation order, the computation of these intermediate results can be interwoven with the computation of the Jacobians, as the Jacobians are computed in program-order. In the left-to-right evaluation there’s no such luck: \( P(x) \) must first be executed, with all intermediate results saved, before the \( J_{\dot{P}}(x) \) may be computed. For this reason, when \( a \approx d \) the right-to-left evaluation order is preferred.

2.0.1 Forward mode AD. Consider the example program \( P \) in Figure 1a. The forward mode AD transformation of \( P \), \( \overline{P} \), is shown in Figure 1b; the transformation involves the core rewrite rule:

\[
\text{let } u = f(a, b) \implies \text{let } v = f(a, b) \text{ and } \dot{v} = \frac{\partial f(a, b)}{\partial a} \dot{a} + \frac{\partial f(a, b)}{\partial b} \dot{b} \tag{2}
\]

This rule generalizes for \( n \)-array functions in the obvious way: for any statement \( \text{let } v = f(a_0, \ldots, a_{n-1}) \), we have \( \dot{v} = \sum_{i=0}^{n-1} \frac{\partial f(a_i, b)}{\partial a_i} \dot{a}_i \).

2
We expose forward mode AD to the user via a function

$$jop : (P : \mathbb{R}^n \rightarrow \mathbb{R}^m) \rightarrow (\overrightarrow{P} : \mathbb{R}^n \rightarrow \mathbb{R}^n \rightarrow \mathbb{R}^m)$$

which, for any differentiable program \( P \) computes \( \overrightarrow{P} \), with the property that \( \overrightarrow{P}(x)(\overrightarrow{x}) = \overrightarrow{J}_P(x) \). In simpler terms, \( \overrightarrow{P} \) takes a point and a direction and returns the derivative of \( P \) at the point and in the given direction by multiplying the direction on the right of the right-to-left association of the Jacobian for \( P \). In order to recover the full Jacobian \( \overrightarrow{J}_P(x) \), \( \overrightarrow{P}(x) \) is mapped over the standard basis of \( \mathbb{R}^n \).

### 2.0.2 Reverse mode AD

The reverse mode AD transformation of \( P, \overrightarrow{P} \), is shown in Figure 1c. As with the forward mode transformation, it involves the core rewrite rule:

$$\begin{align*}
\text{let } v &= f(a, b) \\
\text{let } \overrightarrow{v} &= \overrightarrow{f(a, b)} \overrightarrow{\overrightarrow{v}} \\
\text{let } \overrightarrow{\overrightarrow{v}} &= \overrightarrow{f(a, b)} \overrightarrow{\overrightarrow{\overrightarrow{v}}} \overrightarrow{\overrightarrow{\overrightarrow{\overrightarrow{v}}}}
\end{align*}$$

where \( \overrightarrow{v} \) is the adjoint of \( v \), which is the derivative of \( P \) with respect to \( a \). The adjoint of a variable \( v \) may be updated multiple times—any use of \( v \) may contribute to the derivative (see, e.g., \( x_1 \) in Figure 1c). Reverse mode AD corresponds to computing the left-to-right Jacobian \( \overrightarrow{J}_P(x) \). All intermediate results of \( P \) which appear in adjoint expressions must first be saved, which corresponds to computing \( P \) before the adjoint of each program variable. The \( \overrightarrow{\cdot} \) indicates the presence of these statements, along with any preceding adjoint expressions. The adjoints in Figure 1c are computed in the reverse program order, corresponding to a left-to-right association of the Jacobian of \( P \). Any adjoint \( \overrightarrow{v} \) is necessarily finalized before it is used: all uses of \( v \) must occur after assignment; any contributions to \( \overrightarrow{v} \) must necessarily appear before any uses of \( \overrightarrow{v} \) in reverse program order. \( \overrightarrow{P} \) takes as additional arguments the initial adjoints of the output \( \overrightarrow{\overrightarrow{v}} \) and \( \overrightarrow{\overrightarrow{\overrightarrow{v}}} \) and additionally returns the adjoints of the input, \( \overrightarrow{\overrightarrow{v}} \). Reverse mode AD is made available to the user via a function:

$$vjp : (P : \mathbb{R}^n \rightarrow \mathbb{R}^m) \rightarrow (\overrightarrow{P} : \mathbb{R}^n \rightarrow \mathbb{R}^n \rightarrow \mathbb{R}^m)$$

which computes \( \overrightarrow{P} \) for any differentiable program \( P \), with the property that \( \overrightarrow{P}(x)(\overrightarrow{x}) = \overrightarrow{J}_P(x) \). As \( \overrightarrow{P} \) has the same computational properties as \( J_P(x) \), it is the preferred choice when \( m \ll n \). When \( P \) computes a scalar function (i.e., \( m = 1 \)), \( vjp \) returns the complete gradient of \( P \) in a single pass; \( jop \) requires \( n \) passes, one for each basis vector of \( \mathbb{R}^n \). Programs with very high-dimensional inputs and low-dimensional outputs appear frequently in the fields of optimization and machine learning, making \( vjp \) the computationally superior choice. The \( vjp \) transformation is also more complex due to computing adjoints in reverse order, storing of intermediate program variables, and accumulation of derivatives into adjoint variables, which turn any read of a variable in the original program into a write in the transformed program. For these reasons, we concentrate our discussion on reverse mode AD.

Higher-order derivatives are supported by nesting of \( vjp \) or \( jop \).

### 2.1 Language

We perform our transformation on a data-parallel language which features arbitrary nesting of second-order array combinators (SOACs), e.g., map, reduce, and scan. SOACs are variadic in their number of arguments and returns, i.e., zips and unzips are implicit. For example,

$$\text{unzip3 } (\lambda(a, b, c) \rightarrow (c, b, a)) \space \text{zip3 as bs cs)}$$

may be equivalently written as simply

$$\text{map } (\lambda(a, b, c) \rightarrow (c, b, a)) \space \text{as bs cs}$$

The source language supports higher-order functions, polymorphism, modules, and similar high-level features, which are compiled away using a variety of techniques [12, 20] before we perform AD. The only remaining second-order functions are the SOACs. Lambdas can only appear syntactically in SOACs and VJP/JVP, and are not values. As such we do not suffer from “perturbation confusion” [26]. Further, a significant battery of standard optimisations (CSE, constant folding, aggressive inlining) is also applied prior to AD.

The language is written in A-normal form [39] (ANF); all subexpressions are variables names or constants with the exception of the body expression of loops, if-expressions and let-expressions. let-expressions consist of a series of bindings—that we also call statements—followed by a sequence of one or more returns:

$$\text{let } a = 5 \space \text{let } b = a \space \text{in } b$$

The language is purely functional: re-declarations of the same variable (in a given scope) should be understood as a notational convenience for variable shadowing. It supports a functional flavor of in-place updates based on uniqueness types [18]. The binding let \( xs[i] = x \) is syntactic sugar for let \( xs' = xs \space \text{with} \space [i] \leftarrow x \), which has the semantics that \( xs' \) is a copy of \( xs \) in which the element at index \( i \) is updated to \( x \), but also provides the operational guarantee that the update will be realized in place.

The language also features sequential and pure loops, which have the semantics of a tail-recursive function:

$$\text{let } y = \text{loop } (x) = (x_0) \space \text{for } i = 0 \ldots n - 1 \space \text{do } c$$

The loop is initialized by binding \( x_0 \) to \( x \). Each iteration of the loop executes \( c \), and binds the result of the expression to \( x \), which is used on the subsequent iteration of the loop.
4.1 The Gist of Our Technique

We first demonstrate the gist of our technique on a perfect-nest example in Section 4.1, then present the code transformation across scopes in a semi-formal manner in Section 4.2, and conclude with an example demonstrating our code generation for loops and the time-space trade-off in Section 4.3.

4.2 Transformation Rules Across Scopes

Figure 3 sketches the reverse-mode AD code transformation at the lexical levels of a body of statements $(\omega | p_{body})$, a sequence of statements $(\omega | p_{stmts})$ and individual statements $(\omega | p_{stmt})$. We recall that body refers to a sequence of let bindings (statements) followed by an in result. Our presentation sacrifices some formalism for readability, for example, by omitting the environment, which essentially carries out a mapping from the variables of the source program to their

\[ f(x_1, \ldots, x_n) = [f(x_1, \ldots, x_n)] \]

\footnotetext{Map independently applies a function argument to each of the elements of an array (argument), i.e., map $f \ [x_1, \ldots, x_n] = [f(x_1, \ldots, x_n)]$.}
Rule $vjp_{\text{body}}$ refers to a body $\text{body} = \text{stmts in res}$:

$$vjp_{\text{body}}(\text{res}, \text{stmts in res} \rightarrow \text{stmts in (res, } FV(\text{body})\text{)}$$

where $\text{stmts} \leftarrow vjp_{\text{stmts}}(\text{stmts})$ and $FV(\text{body}) \leftarrow FV(\text{body})$

Rule $vjp_{\lambda}$ refers to a lambda function $\lambda x_1, \ldots x_n \rightarrow \text{body}$:

$$vjp_{\lambda}(\text{R}(\text{res}, \lambda x_1, \ldots x_n \rightarrow \text{stmts in res} \rightarrow \lambda x_1, \ldots x_n \rightarrow \text{body})$$

where $\text{stmts in (res, } FV(\text{body})\text{)} \rightarrow \text{stmts in (res, } FV(\text{body})\text{)}$

body $\leftarrow \text{stmts in } FV(\text{body})$

Rule $vjp_{\text{stmts}}$ simply folds over each statement:

$$vjp_{\text{stmts}}(\text{stmt, stmts} \rightarrow \text{stmts, vjp}_{\text{stmts}}(\text{stmts}), \text{stmts})$$

where $\text{stmts} \leftarrow vjp_{\text{stmts}}(\text{stmt})$

Rule $vjp_{\text{stmt}}$ for scalar multiplication (simple statement):

$$vjp_{\text{stmt}}(\text{let } x = a \cdot b \rightarrow \text{stmts, stmts})$$

where $\text{stmts} \leftarrow \text{let } x = a \cdot b$

$$\text{stmts} \leftarrow \text{let } x = a \cdot b$$

Rule $vjp_{\text{stmt}}$ for loop statement: checkpoints and re-executes body

(Apply only to loops whose variants do not change the shape within the loop)

$vjp_{\text{stmt}}(\text{let } y = \text{loop } (x) = (x_0) \text{ for } i = 0 \ldots n-1 \text{ do } \text{body} \rightarrow \text{stmts, stmts})$

where $\text{stmts} \leftarrow \text{let } x = a \cdot b$

$$\text{stmts} \leftarrow \text{let } x = a \cdot b$$

Figure 3. Reverse AD transformation rules for body of statements, scalar multiplication and do loops.

The $vjp_{\text{stmts}}$ rule highlights the redundant-execution mechanism that removes the need to implement the tape as a separate abstraction: each statement $\text{stmt}$ is processed individually (by $vjp_{\text{stmt}}$), producing a sequence of statements on the forward sweep, denoted $\text{stmts}$, that brings into scope whatever information is necessary to execute the return sweep for that statement, denoted by $\text{stmts}^\ast$.

For example, the forward sweep of a multiplication statement $\text{let } x = a \cdot b$, is the statement itself. Re-executing it brings into scope the value of variable $x$ which may be needed in the return sweep of a following statement, e.g., $\text{let } y = x \cdot a$, which would require an update to the adjoint of $a$ according to the rewrite rule of Equation (3): $a \leftarrow \frac{\partial (x \cdot a)}{\partial a} + \frac{\partial (y)}{\partial a}$, which results in $\overline{a} \leftarrow x \cdot \overline{y}$.

The rule for a loop statement is more complex. Figure 3 shows a loop that exhibits a loop-variant variable $x$—whose value changes through the loop—which is initialized with $x_0$ just before the loop is started, and whose value for the next iteration is bound to the result of the loop body. The result of the entire loop is bound to variable $y$. The rule assumes that the shape of $x$ does not change throughout the loop.

The forward sweep $\text{loop}$ is the original loop, except that its body is modified to checkpoint into array $\text{xs}$ the value of $x$ at the entry of each iteration. As such $\text{xs}$ is also declared as loop variant and initialized to $x_0$, which, in turn, is allocated (by $\text{scratch}$) just before the loop statement. Note that only the loops of the current scope are checkpointed; an inner-nested loop would be re-executed but not checkpointed.

The return sweep consists mainly of a loop that iterates with $i$ from $n-1$ down to $0$: the first statement re-installs the value of $x$ for the current iteration from the checkpoint (i.e., $\text{xs}[i]$). The remaining statements, $\text{stmts}^\ast$, are those generated by $vjp_{\text{body}}$ from the original loop body, including its forward sweep. As $\text{stmts}^\ast$ may use $x$, re-installment is necessary. Re-execution of the forward sweep brings into scope all variables that might possibly be used by the return sweep of the body.

The result of a reversed iteration is the adjoint of the original result $\overline{y}$, together with the (updated) adjoints of the free variables used in the loop $\overline{f_{\text{osdy}}}$, which are declared as variants through the loop $\overline{f_{\text{osdy}}}$. These are declared as variants through the loop $\overline{f_{\text{osdy}}}$, such that the updates of all iterations are recorded. $\overline{f_{\text{osdy}}}$ is the adjoint of the free variables prior to entering the loop; if a free variable was used in a prior statement of the same scope (and hence has an existing potentially non-zero adjoint) its adjoint is simply retrieved. Otherwise, its adjoint is initialized to a zero element of the appropriate type and shape.

---

The statements of the forward and return sweeps are arranged symmetrically; all statements of the forward sweep come before any of the return sweep, and the latter is organized in the reverse order of the original statements: $vjp_{\text{stmts}}(\text{stmt, stmts} \rightarrow \text{stmts, vjp}_{\text{stmts}}(\text{stmts}, \text{stmts})$. 

respective adjoints. Instead, we assume that the adjoint $\overline{x}$ of a variable $x$ is always available.

The $vjp_{\text{body}}$ rule refers to transforming a body of statements, which defines a new scope. The rule starts by binding the body result to its adjoint $\text{res} \rightarrow \overline{\text{res}}$ (not shown). This is safe because the transformation works backwards, hence $\overline{\text{res}}$ is already available from the outer scope (ultimately passed as parameter to $vjp_{\text{function}}$ by the user). The statements of the transformed body $\text{stmts}$ are those generated by $vjp_{\text{stmts}}$ and the result consists of the original result $\text{res}$, extended with the adjoints $\overline{f_{\text{osdy}}}$, consisting of all free variables that are used inside the body. $FV(\text{body})$ finds the free variables used in the body and $FV(\text{body})$ refers to their adjoints.

Rule $vjp_{\lambda}$ refers to an unnamed (lambda) function, and its reverse-AD code is essentially obtained by calling $vjp_{\text{body}}$ on lambda’s body. Note that $x_1, \ldots, n$ are free variables in $\text{body}$, and as such their adjoints are among $\overline{f_{\text{osdy}}}$.
When generating code for the return sweep of a statement with the (adjoint) result \( y \) consists of a loop surrounded by statements in the same parallel constructs do not transformation (3) applied to code containing one loop (1). The figure shows the result of the reverse AD transformation (3) applied to code containing one loop (1). One can notice that the original-loop statements are re-executed twice (\( \text{stms}_\text{loop} \) and \( \text{stms}_\text{loop} \)).

The final statement of the return sweep is \( \text{stms}_\text{x}\); it semantically updates the adjoint of the loop-variant initializer \( x_0 \) with the (adjoint) result \( x' \) of the return sweep loop—the code is generated by \( \text{vp}_{\text{stms}}(\text{let } x' = x_0) \).

Sequential loops are the only construct that require iteration checkpointing: parallel constructs do not (e.g., because map iterations are independent by definition).

For completeness, we conclude by treating array indexing. When generating code for the return sweep of a statement \( \text{let } y = a[i] \), we must update \( a[i] \) with the “contribution” of \( y \). This leads us to the following rule:

\[
\text{vp}_{\text{stms}}(\text{let } y = a[i]) = (\text{let } y = a[i], \text{let } y = \text{upd}_i y a)
\]

The \text{upd} construct merits further elaboration. Semantically, \text{upd}_i a returns \( a \) but with the value at index \( i \) changed to be \( v + a[i] \). Operationally, the array \( a \) is directly modified in-place. To preserve purely functional semantics, we require that the “old” \( a \) and its aliases are never accessed again, similar to the in-place updates of Section 2.1.

4.3 Demonstrating Loops by Example & Tradeoff

Figure 4, Listing 3, demonstrates the result of the \( \text{vp}_{\text{stms}} \) transformation for the source code shown in Listing 1, which consists of a loop surrounded by statements in the same scope (\( \text{stms}_\text{out}, \text{stms}_\text{after} \)). With our strategy, the original loop is executed twice: once on the forward sweep when it is checkpointed (\( \text{stms}_\text{loop} \)), and a second time on the return sweep, where the forward sweep of the loop body (\( \text{stms}_\text{loop} \)) is responsible to bring into scope all the (original) variables computed in the corresponding iteration. If the loop body consists of scalar computations, re-computation is cheap. If the loop body consists of nested recurrences, e.g., other loops, then the re-computation overhead can be optimized, as before, by creating perfect loop nests and flattening them.

The latter is necessary because otherwise, the need of checkpointing will keep them alive, and comes at the cost of (asymptotically) increasing the memory footprint. The time-space tradeoff [42] is demonstrated in Listing 2: applying the reverse AD to the strip-mined loop \( (k = \log_m m^k \text{ times}) \)

5 Rewrite Rules for Parallel Constructs

Having defined the transformation across scopes, this section focuses on the mapping of individual parallel constructs within a scope, though we allow arbitrary nesting of them and loops together. We organize the discussion in free-writing style that focuses on the reasoning that led to the derivation of the rules rather than presenting them in a formal, but dry and verbose notation. Some of these rules resemble some that have been published previously [34], but we developed them independently [30].

5.1 Reduce and Multi-Reduce

We recall that the semantics of reduce, for an arbitrary binary associative operator \( \circ \) and its neutral element \( e_0 \) is:

\[
\text{reduce } \circ e_0 \circ \left\langle a_0, a_1, \ldots, a_{n-1} \right\rangle = a_0 \circ a_1 \circ \ldots \circ a_{n-1}
\]

If the result of \( \text{reduce} \) is let-bound to variable \( y \), we can reason more easily about the contribution of the reduce statement to the adjoint of \( a_i \) if we group terms as:

\[
\text{let } y = (a_0 \circ \ldots \circ a_{i-1}) \circ a_i \circ (a_{i+1} \circ \ldots \circ a_{n-1})
\]

If we would know, for every \( i \), the terms \( l_i = a_0 \circ \ldots \circ a_{i-1} \) and \( r_i = a_{i+1} \circ \ldots \circ a_{n-1} \), we could directly apply the main rule for reverse AD given in Equation (3), which results in:

\[
\frac{\partial l_i}{\partial a_i} \circ a_i \circ r_i \circ y
\]

where \( l_i \) and \( r_i \) are considered constants and \( \mathbb{T} \) denotes a potentially vectorized addition that matches the datatype.
The code for the right-hand side (RHS) can be generically generated as a function \( f \) that can be mapped to all \( a_i, l_i, r_i \):
\[
f \leftarrow \text{\textit{adj}}, \lambda(l_i, a_i, r_i) \rightarrow l_i \odot a_i \odot r_i
\]
extcept that the adjoints of \( l_i \) and \( r_i \) are not returned by \( f \).

Finally, all the \( l_i \) and \( r_i \) values can be computed by a forward and reverse exclusive scan (prefix sum), respectively. Essentially, the forward sweep is the reduce statement. Denoting by \( a_s = [a_0, a_1, \ldots, a_{n-1}] \), and assuming for simplicity that \( \odot \) has no free variables, the return sweep is:
\[
\text{let } is = \text{scan}^{\text{exc}} \odot e_\odot \text{ as } \\
\text{let } rs = \text{reverse as } \rightarrow \\
\text{scan}^{\text{exc}}(\lambda x y \rightarrow y \odot x) e_\odot \rightarrow \text{reverse}
\]
\[
\text{let } as = \text{map } f \text{ as } rs
\]

Essentially, the reverse AD code for an arbitrary reduce requires a map, two scan\(^{10}\) and two reverse operations, which preserve the work-depth parallel asymptotics of the original program. In practice, this is quite expensive,\(^{11}\) but luckily most standard operators, admit more efficient translations:

\subsection{Special Cases of Reduce Operators.}
The treatment of plus, min, max is known [21], but we present a more efficient rule for multiplication. We recount here for completeness and because they are used for reduce-by-index, which, to our knowledge, has not been covered before.

When the reduce operator is (vectorized) plus, we have
\[
\frac{\partial l_i + a_i + r_i}{\partial a_i} = \gamma = \gamma,
\]
hence the return sweep becomes
\[
\text{let } as = \text{map } f \text{ as } rs
\]
This is also derived automatically by the simplification engine from the general rule.

When the reduction operator is multiplication, we have
\[
\frac{\partial (l_i \ast a_i \ast r_i)}{\partial a_i} = l_i \ast r_i \ast \gamma = \gamma,
\]
We discriminate three cases here:

- if all \( a_i \) elements are nonzeros, then \( l_i \ast r_i = \frac{\partial}{\partial a} \gamma \) and 
  \[ y \neq 0, \text{ hence we update each element: } \frac{\partial \gamma}{\partial a}, + = \frac{\partial}{\partial a} \gamma, \]

- if exactly one element at index \( i_0 \) is zero, then \( l_i \ast r_i \) is zero for all other elements, and 
  \[ \frac{\partial \gamma}{\partial a_{i_0}} + = y \ast \gamma \]

- otherwise, if \( \forall i, l_i \ast r_i \equiv 0 \), and \( \gamma \) remains unchanged.

The forward sweep is modified to compute the number of zeros in \( a_i \) and the product of non-zero elements (by a map-reduce operation), followed by setting the reduced result \( y \) accordingly. The return sweep computes the contributions by a parallel map and updates adjoints as discussed before.

Finally, when the reduction operator is \( \text{min} \) (or \( \text{max} \)), then only the adjoint of (one of) the minimal array element should receive the contribution of \( y \), because it is the only one that was used to compute the result. As such, the forward sweep is modified to compute the minimal element together with its (first) index \( i_y \) (the common “argmin” operation), which can be done with a parallel reduction. The

\[ \text{min} = \arg \min \]

forward sweep is then \( \text{let } (y, i_y) = \text{argmin } as \), and the return sweep could be \( \text{let } as[i_y] = y \).

However, at this point we add a compiler improvement that exploits sparsity: For example, if \( as \) has not been created yet, then instead of inserting the update statement, we (statically) record in \( vjp \)'s environment that \( as \) is sparse (currently one non-zero element). Further operators such as reduc condemn, may refine the sparse structure of \( as \) by adding more points to it. If \( as \) is created by a map operation, and \( as \) is still sparse, then we (statically) replace the adjoint code of the map with (only) the adjoint code of the iterations corresponding to the non-zeros in \( as \)—because the other iterations have no effect.

\subsection{Reduce-by-Index.}
Reduce-by-index \([17]\), a.k.a. multi-reduce, essentially generalizes a histogram computation by allowing the values from an array \( (a) \) that fall into the same bin (index from \( \text{inds} \)) to be reduced with an arbitrary associative and commutative operator \( \odot \) having neutral element \( e_\odot \), where the number of bins \( m \) is typically assumed smaller than that of index-value pairs \( n \), i.e.,
\[
\text{let } hs = \text{reduce-by-index } (\odot) e_\odot \text{inds as } \]
has the semantics:
\[
\text{loop } hs = \text{replicate } m e_\odot \text{ for } i = 0 \ldots n-1 \text{ do }
\]
\[
\text{let } hs[\text{inds}[i]] \equiv a[i] \text{ in } hs
\]
A similar reasoning as for the arbitrary reduce suggests that two scans need to be applied to each subset of elements that fall in the same bin, a.k.a., multi-scan, in order to compute the \( l_i \) and \( r_i \) terms for every \( i \). Then the contributions to the adjoint of \( as \) of a reduce-by-index statement are computed as before by \( \text{map } f \text{ as } rs \). Assuming a constant key size, the multi-scan can be implemented within the right work-depth asymptotic by (radix) sorting \( a \) according to the corresponding bins, and then by applying irregular-segmented scans (forward and reverse) on the result. Work is in progress to implement this case. We have implemented however the special-case operators discussed for reduce. Essentially, the forward sweep consists of the reduce-by-index statement, but enhanced with the extended operators, and the return sweep is similar to reduce, except that in the update formula of the adjoint \( as[i] \), \( \gamma \) is replaced with \( hs[\text{inds}[i]] \).

\subsection{Scan or Prefix Sum}
An inclusive scan \([5]\) computes all prefixes of an array by means of an associative operator \( \odot \) with neutral element \( e_\odot \):
\[
\text{scan } \odot e_\odot [a_0, a_{n-1}] \equiv [a_0, a_0 \odot a_1, a_0 \odot a_1 \odot \ldots \odot a_{n-1}]
\]
While the derivation of (multi-) reduce builds on a functional-like high-level reasoning, in scan’s case, we found it easier to reason in an imperative, low-level fashion. For simplicity we assume first that \( \odot \) operates on reals, and generalize later:
The loop above that implements scan, writes each element of the result array \( r[s] \) exactly once. To generate its return sweep, we can reason that we can fully unroll the loop, then apply the main rewrite-rule from Equation (3) to each statement and finally gather them back into the loop below:

\[
\text{let } (\bar{r}, \bar{s}) = \text{scan} (\bar{r}^{-}, \bar{s}^{-}) \quad \text{(copy } \bar{r}^{-}, \bar{s}^{-}) \text{ for } i = n-1 \ldots 0 \text{ do}
\]

\[
\begin{align*}
\bar{r}[i] &\leftarrow \frac{\partial (r[s][i] \odot r[s][i])}{\partial r[s][i]} \odot \bar{s}[i] \quad \text{in } (\bar{r}, \bar{s}) \\
\bar{s}[i] &\leftarrow \frac{\partial (r[s][i] \odot r[s][i])}{\partial r[s][i]} \odot \bar{r}[i]
\end{align*}
\]

Simple dependence analysis, for example based on direction vectors, shows that the loop can be safely distributed across its two statements, since they are not in a dependency cycle:

\[
\begin{align*}
\text{let } (\bar{r}'^{-}) &\leftarrow \text{map } (x \rightarrow \bar{r}[x]) \quad \text{for } i = n-1 \ldots 0 \\
\text{let } \bar{s} &\leftarrow \text{map } (x \rightarrow \bar{s}[x]) \quad \text{in } (\bar{r}', \bar{s})
\end{align*}
\]

The second loop (computing \( \bar{s}^{-} \)) is essentially a map once we know the values of \( \bar{r}'^{-} \). Denoting by \( c_{n-1} = 1 \) and \( c_i = \frac{\partial (r[s][i] \odot r[s][i])}{\partial r[s][i]} \), the first loop (computing \( \bar{r}'^{-} \)) is a backward linear recurrence of the form

\[
\bar{r}'_{n-1} = \bar{r}'_{n-1} -1 \\
\bar{r}'_i = \bar{r}'_i + c_i \cdot \bar{r}'_{i+1}, \quad i = n-2 \ldots 0
\]

where \( \bar{r}' \) is the adjoint of the result of the original loop statement, which is known before the reversed loop is entered.

Such a recurrence is known to be solved with a scan whose operator is linear-function composition [6]. To summarize: the forward sweep is the original scan. The return sweep consists of (1) the map that computes the \( c_i \) values, (2) the scan that computes the backward linear recurrence, and (3) the map that computes that updates \( \bar{s}^{-} \):

\[
\begin{align*}
\text{let } (d, c) &\leftarrow \text{map } (x \rightarrow \text{if } i == n \text{ then } (0, 1) \text{ else } (\bar{r}'[x], c[x]) \text{ for } i = n \ldots 1) \\
\text{let } \bar{s} &\leftarrow \text{scan } 1 n_{\alpha} (d, c) \\
\text{let } \bar{r}' &\leftarrow \text{scan } 1 n_{\alpha} (\bar{r}, (\bar{r}, \bar{s})) \quad \text{(reverse } ds \text{)} \quad \text{(reverse } cs) \\
\text{let } \bar{s}^{-} &\leftarrow \text{map } (x \rightarrow \text{if } i == 0 \text{ then } \bar{r}'[x] \text{ else } \frac{\partial (r[s][i] \odot r[s][i])}{\partial r[s][i]} \odot \bar{r}'[i]) \quad \text{as } [0 \ldots n-1]
\end{align*}
\]

For generalization, we observe that any element type can be linearized to a vector, say size of \( d \). Then we observe that

- A term like \( \frac{\partial (r[s][i] \odot r[s][i])}{\partial a_i} \) corresponds to the Jacobian of the function \( \text{.rt}(y) = r[s][i] \odot y \) at point \( a_i \), denoted by \( f_{\text{rt}}(a_i) \in \mathbb{R}^{d \times d} \).
- A term like \( \frac{\partial (r[s][i] \odot r[s][i])}{\partial r[s][i]} \odot \bar{s}[i] \in \mathbb{R}^d \) applies the Jacobian to a vector and corresponds to \( v_j p_{\lambda}(\bar{s}[i], \text{rt}) \).

It follows that in the rewrite rule above, \( \times \) and \( \odot \) are generalized from scalar multiplication to matrix-vector and matrix-matrix multiplication, respectively, and \( + \) to vector addition. In particular, the \( \alpha_i \) operator of scan, has neutral element \((0 \in \mathbb{R}^d, I \in \mathbb{R}^{d \times d})\), where \( I \) is the identity matrix, and \( \alpha_i \) computes, among others, one matrix multiplication. If \( d \) is a constant, e.g., tuples of scalars, the work-depth asymptotic of the original program is preserved. Otherwise, this rule provides no such guarantee. We have implemented the case when scan's element type is one scalar. We also support vectorized such operators by turning them beforehand to a regular-segmented scan, by the rule:

\[
\text{scan } (\text{map } (\odot)) \bar{r}^{-} \bar{s} \Rightarrow \\
\text{transpose } \bar{s} \circ \text{map } (\text{scan } \circ \odot e_i) \circ \text{transpose}
\]

We treat separately the case of scan with (vectorized) plus operator, in which the contributions to be accumulated by \( \bar{s} \) are given by \( \text{scan } (\odot) 0 \circ \text{reverse } \bar{r}^{-} \circ \text{reverse} \).

### 5.3 Parallel Scatter

The statement \( \text{let } ys = \text{scatter } xs is vs \) produces an array \( ys \) by updating in-place the array \( xs \) (which is consumed) at the \( m \) indices specified in array \( vs \) with corresponding values taken from \( vs \). Scatter has constant depth, and work proportional with \( m \), the size of array \( vs \), i.e., it does not depend on the length \( n \) of the updated array \( ys \). Our rule assumes that \( vs \) contains no duplicates, i.e., idempotent updates are currently not supported.

Denoting by \( \text{gather} \) the operation that reads from a support array elements at a given subset of indices, i.e.,

\[
\text{let } gather \text{ arr inds } = \text{map } (\lambda i \rightarrow xs[i]) \text{ inds}
\]

we forward sweep saves (in \( xs_{\text{saved}} \)) the elements of \( xs \) that are about to be overwritten prior to performing the update:

\[
\text{let } xs_{\text{saved}} = \text{gather } xs
\]

\[
\text{let } ys = \text{scatter } xs is vs
\]

Since the original statement performs \( ys[\text{is}[i]] = vs[i] \) for any \( i \in \text{is} \), we have by Equation (3) \( \text{is}[i] = vs[i] \).

As such, the return sweep (1) first updates the adjoint of \( vs \), then (2) creates the adjoint of \( vs \) by zeroing out the elements from \( \bar{r}^{-} \) that were subject to the scattered update—because those adjoints correspond to elements that were never in \( xs \), and finally, (3) restores \( xs \) to its state before the update:

\[
\text{let } \bar{r}^{-} = \text{gather } \bar{s}^{-}
\]

\[
\text{let } \bar{s}^{-} = \text{scatter } \bar{r}^{-} \bar{s}^{-} (\text{replicate } m) 0
\]

\[
\text{let } xs = \text{scatter } ys is xs_{\text{saved}}
\]

Note that both forward and return sweep preserve the original work-depth asymptotics, because all operations have work proportional to \( m \) (not \( n \)). This would not hold if, e.g., the forward sweep would make a copy of the whole \( xs \).

### 5.4 Map

A map applies a lambda function to each element of an array, producing an array of same length:

\[
\text{let } xs = \text{map } (\lambda x \rightarrow \text{stms in } x) as
\]
If the lambda has no free variables, the return sweep is:

\[
\text{let } \overline{a}S = \text{map}(\lambda(a, \overline{a}, x) \rightarrow \overline{stms} \leftarrow \overline{stms} \text{ in } \overline{a}_0 + \overline{a}) \quad \text{as } \overline{a}S \overline{xs}
\]

where \(\overline{stms}\) and \(\overline{stms}\) denote the forward and reverse statements of lambda’s body, and \(\overline{a}\) is shadowed by \(\overline{stms}\).

A naive way of handling free variables is to turn them into bound variables. E.g. converting \(\text{map}(\lambda(i \rightarrow as[i])\) is into \(\text{map}(\lambda(i, as') \rightarrow as'[i])\) is \((\text{replicate } n as)\) where \(n\) is the size of \(is\). This is fine for scalars, but asymptotically inefficient for arrays that are only partially used, as here, as the adjoint will be mostly zeroes.

In an impure language, we could update the adjoint of a free array variable \(as[i]\) with an operation \(\overline{as}[i]*=p\), implemented with atomics or locks in the parallel case. In our pure setting, we instead introduce accumulators. An array can be “temporarily” turned into an accumulator with \(\text{withacc}\):

\[
\text{withacc } : [d]a \rightarrow (\text{acc}(a) \rightarrow \text{acc}(a)) \rightarrow [d]a
\]

Intuitively we view an accumulator as a “write-only” view of an array. Semantically, accumulators are lists of index/value pairs, each denoting an update of an array. When we use \(\text{upd}\) on an accumulator, we prepend index/value pair to this list, returning a new accumulator. Operationally, \(\text{upd}\) on an accumulator is implemented by immediately updating the underlying array, not by actually maintaining a list of updates in memory (although such an implementation would be semantically valid). The purpose of accumulators is to allow the compiler to continue to reason purely functionally, in particular that all data dependencies are explicit, while allowing efficient code generation based on in-place updates. Our accumulators are similar to generalized reductions [24] or the accumulation effects in Dex [33] and have the same underlying motivation. The main difference is that in Dex, they are an effect, which requires every part of the compiler to be effect-aware.

Free array-typed variables in \(\text{map}\) are thus turned into accumulators while generating return sweep code for the \(\text{map}\), during which we can perform the updates directly. We allow implicit conversion between accumulators and arrays of accumulators, as this allows us to directly \(\text{map}\) them. E.g.

\[
\text{let } xs = \text{map}(\lambda i \rightarrow as[i])\]

results in the return sweep code

\[
\text{let } \overline{xs} = \text{withacc } \overline{xs} (\lambda \overline{xs} \rightarrow \text{map}(\lambda(i, \overline{x}, xs) \rightarrow \text{upd}(i, \overline{x}, xs)) \rightarrow \overline{xs} \overline{xs})
\]

where we treat \(\overline{xs}\) as an array of accumulators when passed to \(\text{map}\), and treat the result of the \(\text{map}\) as a single accumulator. This is efficient because accumulators have no run-time representation, and saves us from tedious boilerplate.

During the lifetime of the accumulator, the underlying array may not be used—this prevents observation of intermediate state. These rules can be encoded in a linear type system and mechanically checked, which we do in our implementation, but exclude from the paper for simplicity.

Accumulators are sufficient to express the adjoint computation inside maps because (1) any read from an array \(d[i]\) is turned into an accumulation on \(\overline{d}[i]\) as discussed at the end of Section 4.2, and (2) the only place on the return sweep where \(\overline{d}\) can be read outside an accumulation statement is the definition of \(a\), which by definition is the last use of \(\overline{a}\), hence it is safe to turn it back into a normal variable there.

6 Implementation and Optimizations

We have implemented the reverse AD transformation as a pass in a copy of the publicly available Futhark compiler. The presented transformation rules were tuned to preserve fusion opportunities, both with constructs from the statement’s sweep, and across statements. Section 6.2 discusses several omitted issues, namely how to optimize checkpointing for the arrays that are constructed by in-place updates inside loop nests, and how to support while loops, and Section 6.3 presents the limitations of our implementation.

Since accumulators were not supported in the original language, we have implemented them throughout the compiler—for the GPU backends, they ultimately boil down to atomic updates, such as \(\text{atomic add}\) in CUDA. Accumulators however, often result in suboptimal performance, because they access memory in uncoalesced fashion and are subject to conflicts, i.e., threads simultaneously accessing the same location. In this sense, Section 6.1 presents optimizations aimed at turning accumulators into more specialized constructs (e.g., \(\text{map-reduce}\)) that can be better optimized.

6.1 Optimizing Accumulators

We demonstrate our optimizations of accumulator on the matrix-matrix multiplication. The code below assumes \(as \in \mathbb{R}^{m \times q}\) and \(bs \in \mathbb{R}^{p \times n}\) and computes \(cs \in \mathbb{R}^{m \times n}\) by taking the dot product of each (pair of) row of \(as\) and column of \(bs\):

\[
\text{let } xs = \text{map}(\lambda i \rightarrow \text{map}(\lambda j \rightarrow \text{sum}(\text{map}(\ast) as[i, : ] \text{bs}[: , j])) \quad [0..n-1]) \quad [0..m-1]
\]

Differentiating the code above results in the return sweep:

\[
\text{let } \overline{xs} = \text{withacc } \overline{xs} \text{ with } \\
\text{map}(\lambda(i, \overline{x}, xs) \rightarrow \text{upd}(i, \overline{x}, xs)) \rightarrow \overline{xs} \overline{xs})
\]

\[
\text{let } \overline{xs} = \text{withacc } \overline{xs} \text{ with } \\
\text{map}(\lambda(i, \overline{x}, xs) \rightarrow \text{upd}(i, \overline{x}, xs)) \rightarrow \overline{xs} \overline{xs})
\]

\[
\text{let } \overline{bs} = \text{upd}(k, j)(as[i, k]) = bs[k, j]\overline{xs}[i, j]
\]

\[
\text{let } \overline{bs} = \text{upd}(k, j)(as[i, k]) = bs[k, j]\overline{xs}[i, j]
\]

\[
\text{in } (\overline{xs}, \overline{bs})
\]

---

12For simplicity we treat only single-dimensional arrays in this section, but the idea also works in the multi-dimensional case. This type for \(\text{withacc}\) allows only a single result corresponding to the array being updated. In practice, we also need to be able to return an arbitrary secondary result.

13https://github.com/diku-dk/futhark
which is not efficient, because (temporal) locality is sub-utilized. In this sense, we have designed and implemented a pass aimed at turning (common cases of) accumulator accesses into reductions. We present the analysis at an intuitive level: The analysis searches for the first accumulator directly nested in a perfect map-nest and checks whether its indices are invariant across any of the parallel dimensions. In our case as is accumulated on indices \([i,k]\) that are both invariant to the outermost parallel index \(j\). In such a case, the map-nest is split into two: the code on which the accumulated statement depends, and the code without the accumulator statement.\(^{14}\) which is simplified and treated recursively. The map-nest encapsulating the accumulation is reorganized such that the invariant dimension \((j)\) is moved innermost.\(^{15}\) The accumulated statement is taken out of this innermost map, which is modified to produce (only) the accumulated values, whose sum \((\text{reduce} \ (+) \ \emptyset)\) is re-written to be the value accumulated by \(a\):

\[
\begin{align*}
\text{let } (\overleftarrow{\text{a}}) &= \text{withacc}(\overleftarrow{\text{a}}) & & \text{map } (\lambda i \overleftarrow{\text{a}} \rightarrow \overleftarrow{\text{a}}) \\
\text{map } (\lambda k \overleftarrow{\text{a}} \rightarrow \overleftarrow{\text{a}}) & & \text{let } s = \text{sum}(\text{map}(\rightarrow) \text{ bs}[:,j] \overrightarrow{\text{a}}[i,:]) \\
\text{in } & & \overrightarrow{\text{a}}[i,k] += \text{bs}[k,j]\overrightarrow{\text{a}}[i,j] \\
& & \{[0..q-1] \overleftarrow{\text{a}} \} \{[0..n-1] \overleftarrow{\text{a}} \} \\
\text{let } (\overleftarrow{\text{b}}) &= \text{withacc}(\overleftarrow{\text{b}}) & & \text{map } (\lambda k \overleftarrow{\text{b}} \rightarrow \overleftarrow{\text{b}}) \\
\text{map } (\lambda j \overleftarrow{\text{b}} \rightarrow \overleftarrow{\text{b}}) & & \text{let } s = \text{sum}(\text{map}(\rightarrow) \text{ as}[:,j] \overrightarrow{\text{b}}[i,:]) \\
\text{in } & & \overrightarrow{\text{b}}[i,k] += \text{as}[i,k]\overrightarrow{\text{b}}[i,j] \\
& & \{[0..n-1] \overleftarrow{\text{b}} \} \{[0..q-1] \overleftarrow{\text{b}} \}
\end{align*}
\]

One can notice that the code now essentially consists (as expected) of two matrix-multiplication-like kernels. These are optimized by a later pass that performs block and register tiling whenever it finds an innermost map-reduce whose input arrays are invariant to one of the outer-parallel dimensions. We have extended this pass (1) to support accumulators, (2) to keep track of the array layout—i.e., transposed or not, (3) to copy from global to shared memory in coalesced fashion for any layout, and (4) to exploit some of the parallelism of the innermost dot product as inspired from [37].

This optimization is responsible for nearly a one-order-of-magnitude speedup at application level for benchmarks dominated by matrix multiplication such as GMM and LSTM, evaluated in sections 7.6 and 7.7.

6.2 Loop optimizations

As discussed in section 4, by default, loop-variant variables are saved/checkpointed at the entry of each iteration. This technique does not preserve the work asymptotic of the original program when a loop variant array is modified in place. For example, the contrived dependent loop below constructs an array of length \(n\) in \(O(n)\) work, but the checkpointing of the forward sweep would require \(O(n^2)\) work:

\[
\text{loop } (xs) = (xs_0) \text{ for } i = 1..n-1 \text{ do } \\
xs \text{ with } [i,j] = ass[i,j] + xs[i-1,j]
\]

One can observe however that iteration-level checkpointing is not needed if the loop nest does not exhibit any false dependencies (WAR+WAW):\(^{16}\) since no value is “lost” through the loop nest, it is enough to checkpoint \(xs\) only once at the entry to the outermost loop of the nest. Moreover, re-execution is safe because all the over-writes are idempotent. In this sense, we allow the user to annotate such loop parameters that are free of false dependencies: we checkpoint them at the loop-nest entry and restore this copy just before entering the return sweep of that nest. Essentially, we consider that false dependencies are properly named as such, and the user should not rely on them, especially in a data-parallel context. Work in the automatic parallelization area can be used to automatically check the safety of such annotations, statically [16], dynamically [11], and anywhere in between [31].

A second issue relates to while loops, on which we cannot perform AD directly, because their statically-unknown iteration count hinders the allocation of the checkpointing tape. We provide two mechanisms to address this issue: first, the user may annotate a while-loop with an iteration bound \(n\). The while loop is then transformed into an \(n\)-iteration for-loop that contains a perfectly nested if, which only executes the valid iterations of the while loop. Alternatively, in the absence of such annotation, the loop count can be computed dynamically by an inspector—i.e., loop slice that computes only the number of iterations of the loop.

6.3 Current Implementation Limitations

We have already discussed limitations during the presentation of individual constructs. The main remaining one is that we do not currently support loop-variant parameters that change their shape throughout the loop. In principle this can be handled by dynamic re-allocations, albeit this might prove expensive in a GPU setting.

Another shortcoming is that when an in-place update occurs inside an if-then-else, we currently save the target array at the branch’s entry: in principle, we should instead propagate the saved element(s)—denoted \(xs_{saves}\) in Section 5.3—outside branches, and back in to reach the return sweep.\(^{17}\) By construction, this propagation cannot escape the scope of the innermost-enclosing recurrence.

\(^{14}\)The absence of false dependencies, means that the loop has only true (RAW) dependencies or no dependencies at all.

\(^{15}\)It is always safe to interchange parallel loops inwards.
7 Experimental Evaluation

7.1 AD-Bench: Sequential AD Performance
ADBench is a collection of benchmarks aimed at comparing different AD tools [43]. We have used the ADBench framework to measure Futhark versions of the four ADBench problems. We compile to sequential CPU code and report the time taken to compute the full Jacobian relative to the time taken to compute the objective function, using the largest default dataset for each benchmark. We compare against Tapenade [2] and manually differentiated programs. The results are shown in Table 1.

Futhark does well, in particular managing to outperform Tapenade in four out of five cases. For the exception, BA, the bottleneck is packing the result (which is a sparse Jacobian) in the CSR format expected by the tooling, which is code that is not subject to AD. The HAND benchmark has two variants: a “simple” one that computes only the dense part of the Jacobian, and a “complicated” one that also computes a sparse part. Tapenade handles the latter poorly. Both Tapenade and Futhark perform poorly when compared to the hand-written code. Both BA and HAND produce sparse Jacobians where the sparsity structure is known in advance, which is exploited by passing appropriate seed vectors to jvp/vjp.

7.2 Parallel Hardware and Methodology
We benchmark on two different Linux systems: one with two AMD EPYC 7352 CPUs and an NVIDIA A100 GPU running CUDA 11.2, which we denote A100, and another with two Intel E5-2650 CPUs and an NVIDIA RTX 2080Ti GPU running CUDA 11.3, which we denote 2080Ti. We report mean runtime for 10 runs, which includes all overheads, except transferring program input and result arrays between device and host. We report the absolute runtime of the differentiated and primal program, and the “overhead” of differentiation that corresponds to the ratio between the two. In optimal AD, this ratio (counted in number of operations) is supposed to be a small constant [15], hence the ratio serves as a good measure of the efficiency of an AD implementation. Futhark benchmarks using 32-bit floats were performed with block and register tile sizes of 16 and 4, respectively; on 64-bit floats the register tile size is 2.

<table>
<thead>
<tr>
<th>Tool</th>
<th>BA</th>
<th>D-LSTM</th>
<th>GMM</th>
<th>HAND</th>
<th>Comp.</th>
<th>Simple</th>
</tr>
</thead>
<tbody>
<tr>
<td>Futhark</td>
<td>13.0×</td>
<td>3.2×</td>
<td>5.1×</td>
<td>49.8×</td>
<td>45.4×</td>
<td></td>
</tr>
<tr>
<td>Tapenade</td>
<td>10.3×</td>
<td>4.5×</td>
<td>5.4×</td>
<td>3758.7×</td>
<td>59.2×</td>
<td></td>
</tr>
<tr>
<td>Manual</td>
<td>8.6×</td>
<td>6.2×</td>
<td>4.6×</td>
<td>4.6×</td>
<td>4.4×</td>
<td></td>
</tr>
</tbody>
</table>

Table 1. Time to compute the full Jacobian relative to time to compute the objective function. Lower is better.

7.3 Compared to Enzyme
RSBench and XSBench are CUDA implementations of Monte Carlo neutron transport algorithms that are reported in the Enzyme paper [29], and which we have ported to Futhark in order to quantitatively compare our solution with Enzyme. They each constitute a large problem that is suitable for use with forward-mode vjp (the R argument is set to 1). Further, as the Hessian for f has nonzero entries

<table>
<thead>
<tr>
<th>Tool</th>
<th>(k, n, d)</th>
<th>Futhark</th>
<th>AD</th>
<th>PyTorch</th>
</tr>
</thead>
<tbody>
<tr>
<td>RSBench</td>
<td>(5, 494019, 35)</td>
<td>9.3ms</td>
<td>36.6ms</td>
<td>44.9ms</td>
</tr>
<tr>
<td>XSBench</td>
<td>(1024, 10000, 256)</td>
<td>9.9ms</td>
<td>9.6ms</td>
<td>11.2ms</td>
</tr>
</tbody>
</table>

Table 2. Results for RSBench and XSBench. We report the absolute runtimes of the original program and the un-differentiated Futhark program, as well as the overhead of a single forward and return sweep of the reverse-mode differentiated program compared to the un-differentiated one.

7.4 Case Study 1: Dense k-means clustering
k-means clustering is an optimization problem where given n points P in a d-dimensional space we must find k points C that minimize the cost function

$$ f(C) = \sum_{p \in P} \min_{c \in C} ||p - c|| $$

Solving this with Newton’s Method requires computing the Jacobian and Hessian of f. The cost function is easily written with nested map and reduce operations, for which the Hessian can be computed by nesting vjp inside of jvp. Note that while

$$ f : \mathbb{R}^{k \times d} \rightarrow \mathbb{R} $$

and so needs reverse-mode vjp, the differentiated function

$$ vjp f : \mathbb{R}^{k \times d} \rightarrow \mathbb{R} \rightarrow \mathbb{R}^{k \times d} $$

is suitable for use with forward-mode jvp. The results (Table 2) show the overhead caused by our AD and the one reported in the Enzyme paper. Our overhead is slightly smaller, although this may come down to micro-optimisations.
only along the diagonal, we need only a single invocation of \( \text{jvp} \) to compute it, returning exactly this diagonal. This shows how \( \text{jvp/vjp} \) allows the user to exploit sparsity. Further, the nesting of \( \text{jvp/vjp} \) allows us to avoid duplicating the expensive search operation in the cost function, which would not be the case if we were limited to dedicated constructs for Jacobian and Hessian computation.

\( k \)-means clustering can also be solved using a method based on repeatedly calculating histograms \cite{17}, which can be implemented using Futhark’s dedicated histogram construct. The code that is generated by AD consists of maps that perform semantically equivalent histogram-like accumulator updates. Essentially, using AD leads to basically the same code one would write by hand. The main difference is that in the hand-written code, some of the updates have are manually sequentialized, while the AD version maximizes parallelism to degree that is detrimental for some datasets.

In PyTorch we must take care when computing the all-pairs Euclidean distance from data points to the centers. Expressing this computation with broadcasting suffers from massive memory overhead. We get around this by explicitly distributing the quadratic terms in the Euclidean distance. The AutoGrad module computes the Jacobian and Hessian, instead of in one go as we can when nesting \( \text{jvp/vjp} \).

We benchmark with two qualitatively different datasets, with results in Table 3. When the histograms benefit from the optimisations discussed in \cite{17}, the hand-written implementation has a speedup of 4x over our AD approach, while there is no significant difference if they cannot. Our AD approach is slightly faster than PyTorch (as high as 1.22x on A100 and 1.18x on 2080Ti).

7.5 Case Study 2: Sparse \( k \)-means clustering

We have implemented and tested a sparse formulation of \( k \)-means, that uses a dense representation for centroids and a sparse representation of data. Futhark implementations are relatively straightforward adaptations of the dense \( k \)-means to use the CSR format instead.

The PyTorch implementation uses the COO format because AD (\( \text{jvp and hvp} \)) currently raises runtime errors with CSR format.\footnote{We have an open (unresolved) query on the PyTorch discord asking if CSR will be supported in the future.} To efficiently compute the cost function we expand the all pairs norm between data \( d \) and centroids \( c \):

\[
||d_i - c_j||^2 = d_i^2 + c_j^2 - 2d_i c_j^T. 
\]

In expanded form, all terms can be computed using vectorized operations. Due to the array (tensor) presentation \( d_i^2 \) and \( d_i c_j^T \) are computed using PyTorch specialized sparse module, in particular, the (sparse) matrix product uses \textsc{sparse.mm} which allows sparse gradient computation (but also forces the COO representation).

Table 4 shows the runtime results for A100 on three publicly available NLP workloads for \( k = 10 \). Our AD is slower than the manual code by a factor between 2.5—3.7x because the generalized-histogram implementation of the manual code uses a multi-pass technique that allows the updates to fit in the L2 cache \cite{17}. PyTorch AD is more than 400x slower than our AD, which we do not know how to explain.

7.6 Case Study 3: GMM

To evaluate the parallelism-preservation of our AD transformation, we compile the GMM benchmark from the AD-Bench suite to parallel CUDA. We compare against AD-Bench’s implementation of GMM in PyTorch (also run on CUDA), which we have improved (by a \( \geq 10 \times \) factor) by vectorizing all comprehensions. We benchmark on a selection of 1,000 and 10,000-point datasets from AD-Bench featuring a variety of \( d \) (the dimensionality of the input data) and \( K \) (the number of Gaussian distributions used in the model) values.
see Table 5a. We observed that the runtime of the primal program is dominated by matrix multiplication (~70%).

Matrix multiplication is a primitive in PyTorch [32]; we reasonably expect that differentiation of matrix multiplication may be readily implemented very efficiently. In Futhark there are no such primitives: matrix multiplication is written with maps, whose differentiation yield accumulators, which are further optimized as described in Section 6.1.

The benchmark results are shown in Tables 5b and 5c for the A100 and 2080Ti systems, respectively. On the A100 64-bit floats were used; on the 2080Ti system 32-bit floats were used due to the limited double-precision performance (Torch’s reason). The lengths are close to the average sentence length for the 64-bit floats were used; on the 2080Ti and A100 systems. As with GMM, LSTM is dominated by matrix-multiplication computations. Neither AD-based implementation is competitive against the manual cuDNN-based implementation—we suspect that torch.nn.LSTM switches the computation to tensor cores (F16) internally, because the speedups over Futhark are well correlated with the ratio between peak F32 and F16 performance of the two systems.

### 8 Related Work

Reverse-mode differentiation of reduce and scan is discussed in [34]. Our rule for reduce is similar, but we handle scan differently. Our approach is less efficient for complex operands because we manifest the Jacobian matrices, but more efficient for single-value operands on GPUs as it requires less shared memory to implement the derived scan operator. Neither our scan rule nor the one from [34] is asymptotically-preserving in the general case, but they are for most scans that occur in practice.

$F$ is a functional array language that supports nested parallelism. Its AD implementation uses the forward mode, along with a handful of rewrite rules that allow it to exploit sparsity in some cases [41].

Dex is a recent language built specifically to support efficient AD. Empirical benchmarks for AD in Dex have not yet been published, but we can compare their approach [33]. In contrast to our conventional "monolithic" approach where reverse-mode AD is a transformation completely distinct from forward-mode, Dex uses a technique where the program is first "linearized", producing a linear map, after which this linear map is then "transposed", producing the adjoint code. Like Dex, we do not support recursion or AD of higher-order functions. Dex does not make direct use of a “tape” in the classical sense, but instead constructs arrays of closures followed by defunctionalization. The actual run-time data structures will conceptually consist of multiple tapes in the form of multidimensional irregular arrays. Dex does not yet make use of checkpointing, or optimization of particular accumulation patterns as in Section 6.1.

Enzyme shows the advantage of performing AD after standard compiler optimisations has simplified the program [28]. Like Enzyme, we also apply our AD transformation on a program that has already been heavily optimized by the compiler. But where Enzyme is motivated by performing AD on a post-optimization low-level representation, our work takes advantage of both pre-AD optimization, as well as the information provided by high-level parallel constructs. Enzyme has also been applied to GPU kernels [29]. We achieve equivalent performance, but our approach is not based on differentiating single kernels—indeed, the GPU code we generate for a differentiated program may have a significantly different structure than the original program. For example, the optimized adjoint code for a matrix multiplication requires two matrix multiplications, each its own kernel, as in the LSTM and GMM benchmarks.

<table>
<thead>
<tr>
<th>Speedups (×)</th>
<th>Overheads (×)</th>
</tr>
</thead>
<tbody>
<tr>
<td>PyT. Jacob.</td>
<td>Fut. cuDNN</td>
</tr>
<tr>
<td>D0</td>
<td>51.9ms</td>
</tr>
<tr>
<td>D1</td>
<td>173.7ms</td>
</tr>
<tr>
<td>D0</td>
<td>3.1</td>
</tr>
<tr>
<td>D1</td>
<td>3.0</td>
</tr>
<tr>
<td>D0</td>
<td>3.1</td>
</tr>
<tr>
<td>D1</td>
<td>3.2</td>
</tr>
<tr>
<td>D0</td>
<td>2.6</td>
</tr>
<tr>
<td>D1</td>
<td>2.5</td>
</tr>
<tr>
<td>D0</td>
<td>2.6</td>
</tr>
<tr>
<td>D1</td>
<td>3.2</td>
</tr>
</tbody>
</table>

Table 6. LSTM measurements on datasets $D_0$ : $(bs, n, d, h) = (1024, 20, 300, 192)$ and $D_1$ : $(bs, n, d, h) = (1024, 300, 80, 256)$. Py.T, Fut., and cuDNN refer to the PyTorch, Futhark, and the cuDNN-based torch.nn.LSTM implementations.
We have presented a fully operational compiler implementation of both reverse and forward mode AD in a nested-parallel functional language. Our experimental evaluation shows that our mapping of parallel construct and our technique of implementing the tape based on redundant computation is practically effective, and competitive with both (1) well-established frameworks that encompass more specialized languages such as PyTorch, and with (2) newer research efforts aimed at a lower-level language, such as Enzyme.

**References**


9 Conclusions

We have presented a fully operational compiler implementation of both reverse and forward mode AD in a nested-parallel functional language. Our experimental evaluation shows that our mapping of parallel construct and our technique of implementing the tape based on redundant computation is practically effective, and competitive with both (1)
Neural Networks and Deep Learning (1st edition, 2016) by Ian Goodfellow, Yoshua Bengio, and Aaron Courville

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