

- $_{\scriptscriptstyle 1}$ $\Phi_{
 m ML}$: A Science-oriented Math and Neural Network
- ² Library for Jax, PyTorch, TensorFlow & NumPy
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Summary

 $\Phi_{\rm ML}$ is a math and neural network library designed for science applications. It enables users to quickly evaluate many network architectures on their data sets, perform (sparse) linear and non-linear optimization, and write differentiable simulations that scale to n dimensions. $\Phi_{\rm ML}$ is compatible with Jax, PyTorch, TensorFlow and NumPy, and user code can be executed on all of these backends. The project is hosted at https://github.com/tum-pbs/PhiML under the MIT license.

Statement of need

Machine learning (ML) has become an essential tool for scientific research. In recent years, ML has been used to make significant advances in a wide range of scientific fields, including chemistry (Butler et al., 2018), materials science (Wei et al., 2019), weather and climate prediction (Bochenek & Ustrnul, 2022; Rolnick et al., 2022), computational fluid dynamics (Brunton et al., 2020), drug discovery (Jumper et al., 2021; Vamathevan et al., 2019), astrophysics (De La Calleja & Fuentes, 2004; Ntampaka et al., 2015; Petroff et al., 2020), geology (Rodriguez-Galiano et al., 2015), and many more. The use of ML for scientific applications is still in its early stages, but it has the potential to revolutionize the way that science is done. ML can help researchers to make new discoveries and insights that were previously impossible.

ML in science sets itself apart from other ML applications by a number of features.

- The dynamics of the observed system is often (partially) known and can be explicitly simulated. Making use of this knowledge has been shown to improve results when training ML models (Raissi et al., 2019; Um et al., 2020).
- Data typically represent objects or signals that exist in space and time. Data dimensions
 are interpretable, e.g. vector components, time series, n-dimensional lattices.
- Information transfer is usually local, resulting in sparsity in the dependency matrix between objects (particles, elements or cells).
- A high numerical accuracy is desirable, often requiring 64-bit floating point calculations.

However, current machine learning frameworks have limited support for these features, or they are cumbersome to use. $\Phi_{\rm ML}$ is a scientific computing library based on Python 3 (Van Rossum & Drake, 2009) that aims to address these issues and simplify scientific code in the process. It consists of a high-level NumPy-like API geared towards writing easy-to-read and scalable simulation code, as well as a neural network API designed to allow users to quickly iterate over many network architectures and hyperparameter settings. Similar to eagerpy (Rauber et al., 2020), $\Phi_{\rm ML}$ integrates with Jax (Bradbury et al., 2018), PyTorch (Paszke et al., 2019), TensorFlow (Abadi et al., 2016) and NumPy (Harris et al., 2020), providing a custom Tensor class. However, unlike eagerpy, $\Phi_{\rm ML}$'s Tensor adds additional functionality to make user code



- 41 more concise and easier to read.
- $\Phi_{
 m ML}$ has been in development since 2019 as part of the $\Phi_{
 m Flow}$ (Holl et al., 2020) project
- 43 where it originated as a unified API for TensorFlow and NumPy, used to run differentiable
- 44 fluid simulations. With $\Phi_{
 m Flow}$ version 2.0 and consecutive releases, $\Phi_{
 m ML}$ underwent a drastic
- overhaul. A major issue with the previous API, and in fact all popular ML APIs, is the need
- for reshaping, which can quickly get out of hand for physical simulations. The work towards
- automatic reshaping sparked most of the changes that have been made to the library since.
- $_{\mbox{\tiny 48}}$ $\,$ We will first explain the design principles underlying $\Phi_{\rm ML}$'s development, before detailing the
- 49 major design decisions and resulting architecture. For a list of supported features, see the
- 50 GitHub homepage.

51 Design Principles

Here, we lay out our goals in developing $\Phi_{
m ML}$, which serve as the foundation for the design.

33 Reusability

- Simulation code based on $\Phi_{
 m ML}$ should be able to run in many settings without modification.
- 55 The dynamics of a system, e.g. governed by a partial differential equations, are often formulated
- 56 in a dimension-agnostic manner. Simulation code implementing these dynamics should also
- 57 exhibit that property. Most simulations use some form of discretization, such as particles or
- 58 grids. Simulation code written for one such discretization should be easy to port to another
- 59 appropriate one.

50 Compatibility

- 161 There are many toolkits and libraries extending ML frameworks with specialized functionality.
- These are generally only available for a certain framework, be it TensorFlow, PyTorch or Jax.
- $_{63}$ $\Phi_{
 m ML}$ users should be free to choose whatever framework they desire without modifying their
- 64 simulation code. Additionally, simulations should be able to run on GPUs and CPUs and be
- vectorizable without modification, $\Phi_{
 m ML}$ should support Linux, Windows and Mac.

66 Usability

- $_{67}$ Φ_{ML} should be easy to learn and use. To achieve this, the API should be intuitive with
- expressively named functions matching existing frameworks where possible. User code as well
- 69 as built-in simulation functionality should be easy to read, i.e. concise and expressive. We give
- 70 a more detailed explanation of easy-to-read code below.

71 Maintainability

- Users should be able to read and understand all high-level source code of $\Phi_{
 m ML}$. All relevant
- 73 framework functions should undergo continuous testing to ensure patches do not break existing
- $_{74}$ code. When installing $\Phi_{
 m ML}$, users should be able to check the installation status and get hints
- ₇₅ as to how to solve potential issues.

76 Performance

- $_{77}$ Code using $\Phi_{
 m ML}$ should be able to make use of hardware accelerators (GPUs, TPUs) where
- 78 possible. During development, we prioritize rapid code iterations over execution speed but the
- 79 completed code should run as fast as if written directly against the chosen ML library.



Major Design Decisions

Support for Jax, PyTorch, TensorFlow & NumPy

A large fraction of scientific code is re-written one or multiple times due to different preferences in programing languages and libraries. To avoid this as much as possible and reach a large audience, we decided to make $\Phi_{\rm ML}$ compatible with all major Python-based ML libraries as well as NumPy, which they all integrate with. To realize this, we employ the adapter pattern (Freeman et al., 2004), creating an abstract Backend class with adapter subclasses for NumPy, TensorFlow, PyTorch and Jax. This API operates directly on backend-specific tensors, and we use it to implement low-level functions, such as linear algebra routines and neighborhood search. However, writing code that actually runs with all backends requires advanced knowledge of all backends due to the subtle differences between them. PyTorch, for example, does not allow negative steps in tensor slices and TensorFlow does not support assigning values to slices.

Custom Tensor class

The differences between the backends motivate us to provide a Tensor class that handles consistently across all backends. It also enables most of the additional functionality described below, making it easier to write reusable code. A $\Phi_{\rm ML}$ tensor wraps and extends a tensor from one of the supported backends. To operate efficiently on $\Phi_{\rm ML}$ tensors, we include a NumPy-like public API while relegating the Backend API to internal use. The public API takes in $\Phi_{\rm ML}$ tenors, determines the appropriate Backend, and calls the corresponding low-level function. Since all backend-specific tensors are represented by the same Tensor class in $\Phi_{\rm ML}$, code written against $\Phi_{\rm ML}$'s public API is backend-agnostic. Data can also be passed between backends, internally using the tensor sharing functionality of DLPack (al., 2017) when possible. This way, an easy-to-use PyTorch network can interact with a Jax simulation for performance but also with an identical PyTorch simulation to facilitate debugging.

Named dimensions

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In $\Phi_{
m ML}$, dimensions are not referenced by their index but by name instead. We make dimension names mandatory for all dimensions, forcing users to explicitly document the meaning of each dimension upon creation. The name information gets preserved by tensor manipulations and can be inspected at any later point, e.g. by printing it or using a debugger. Named dimensions are also present in other numerics libraries, such as pandas (McKinney, 2010), xarray (Hoyer & Hamman, 2017), einops (Rogozhnikov, 2018), and are available for PyTorch as an add-on (NLP, 2019). However, these libraries make dimension names optional and, consequently, cannot support them to the same extent that Φ_{ML} can, preventing mainstream adoption. In $\Phi_{
m MI}$, dimension names are one part of a carefully-designed set of tools, making them more intuitive and useful than in previous libraries. For instance, $\Phi_{
m ML}$ introduces the convenience slicing syntax tensor.dim_name[start:stop:step], replacing the less readable slices tensor[..., start:stop:step, :], and supports dimension names in all functions as first-class citizens. While naming dimensions adds a small amount of additional code, this is easily outweighed by the gains in readability and ease of debugging. Furthermore, dimension names enable automatic reshaping, which eliminates the need for reshaping operations in user code, often significantly reducing the amount of required boilerplate code.

Automatic reshaping

Named dimensions make it possible to perform reshaping, transposing, squeezing and unsqueezing operations completely under-the-hood. $\Phi_{\rm ML}$ realizes this by aligning equally-named dimensions. Take the operation a + b where a has dimensions (x, y) and b has (y, z). Then $\Phi_{\rm ML}$ will expand a by z and b by x so that both arguments have the common shape (x,y,z) before adding them. This automatic reshaping eliminates the vast majority of shape-related errors as user code is agnostic to the dimension order by default.



Element names along dimensions

In addition to naming dimensions, $\Phi_{\rm ML}$ also supports naming slices or *items* along dimensions. This is optional but highly recommended for dimensions that enumerate interpretable quantities, such as vector components (x, y, z). UnifyML can then check at runtime that the component order is consistent, i.e. that no vector (z, y, x) is added to an (x, y, z)-ordered quantity. Additionally, the slicing syntax becomes more readable when using item names, e.g. tensor.vector['x'] instead of the traditional tensor[:, 0, ...] or tensor[:, -1, ...] (PyTorch dimension order).

136 Non-uniform tensors

With some data structures, such as staggered grids, the number of elements along one or multiple dimensions can be variable. We will refer to tensors holding such data as non-uniform tensors, but they are also known as ragged or nested tensors. Users will often pad the missing elements with zeros to make the data easier to handle but this can lead to problems down the line. Instead, $\Phi_{\rm ML}$ automatically creates non-uniform tensors when stacking tensors with non-matching shapes. The shape attribute of a non-uniform tensor stores its exact layout, allowing users to operate on non-uniform shapes like on regular shapes, e.g. allocating new memory with zeros(non_uniform_shape).

145 Unified functional math

For differentiation, just-in-time compilation and iterative solves, we adopt a function-based approach similar to Jax. This is different from TensorFlow, where gradients are tracked via Python context managers, and PyTorch, where gradients are attached to tensors. $\Phi_{\rm ML}$ unifies these different paradigms, providing unified function operations that run with all backends. For example, math.functional_gradient(f) returns a function that computes the gradient of f and, to solve a sparse system of linear equations, users simply supply a Python function and the desired output of that function.

153 Dimension types

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In all backend libraries, tensor operations act only on certain dimensions, determined by the dimension index in the shape. This behaviour is generally not consistent for all backend libraries.

Consider the response to extra leading dimensions in PyTorch:

- Most functions treat leading dimensions as batch dimensions, i.e. they deal with slices independently, either sequentially or in parallel.
- Some functions like bincount do not allow extra dimensions.
- Reduction functions like sum or prod reduce leading dimensions by default.
- Some functions, such as histogram, flatten all input dimensions.
- Some functions, such as pad, only allow a certain number of leading dimensions.

 Φ_{ML} solves these issues by assigning a type to each dimension. Each of the five allowed types, batch, spatial, instance, channel, and dual, determines how math functions act on dimensions of that type. Spatial operations like fft only act on spatial dimensions and all functions accept tensors with any number of batch dimensions which are always preserved in the operation. Importantly, the order of dimensions is irrelevant to all math functions, only the types matter. For an explanation of all dimension types and further advantages of this system, see the online documentation.

170 Floating-point precision by context

Specifying the floating point precision can be a major headache in computing libraries. NumPy automatically up-casts data types (bool \rightarrow int \rightarrow float \rightarrow complex) and floating point precision (16 bit \rightarrow 32 bit \rightarrow 64 bit). This can cause unintentional data type conversions



when trying to run code with a different precision, as new arrays are FP64 by default. To avoid these issues, TensorFlow has completely disabled automatic type conversion and Jax has disabled FP64 by default. $\Phi_{\rm ML}$ solves the data type problem by enabling automatic casting but determining the desired floating point precision from the operation context rather than the data types of its inputs. The precision can be set globally or specified locally via context managers. All operations automatically convert tensors of non-matching data types. This avoids data-type-related problems and errors, as well as making user code more concise and cohesive.

182 Lazy stacking

Simulations often perform component-wise operations separately if there is no function achieving the desired effect with a single call, like computing the x, y and z-component of a velocity 184 field in three lines. This often leads users to declare separate variables for the components to 185 avoid repeated tensor stacking and slicing. However, this clutters the code and prevents it 186 from being dimension-agnostic. Instead, $\Phi_{
m ML}$ performs lazy stacking by default, i.e. memory 187 is only allocated once the stacked data is required as a block. Consequently, functions can 188 unstack the components, operate on them individually, and restack them, without worrying 189 about unnecessary memory allocations. This system also facilitates stacking tracer tensors, 190 which cannot be done eagerly.

192 Just-in-time compilation

While the previous features allow for concise, expressive and flexible code, the added abstraction layer and shape tracking induces an additional performance overhead. To avoid this in production, $\Phi_{\rm ML}$ supports just-in-time (JIT) compilation for PyTorch, TensorFlow and Jax. Once compiled, only the tensor operations are executed, eliminating all Python-based overhead.

Sparse matrices from linear functions

Solving linear systems of equations is a key requirement in both particle and grid-based 198 simulations. Since the physical influence is typically limited to neighboring sample points or 199 particles, the resulting linear systems are often sparse. Constructing such sparse matrices by 200 hand yields code that is hard to understand and debug as well as limited to specific boundary conditions. Instead, Φ_{ML} lets users specify linear systems with a linear Python function, like with matrix-free solvers. However, these functions often consist of many individual operations, 203 which makes it inefficient to call them at each solver iteration. To avoid this overhead, $\Phi_{
m ML}$ can convert most linear and affine functions to sparse matrices so that solvers can perform the matrix multiplication in a single operation. When JIT-compiling a simulation that includes a 206 linear solve, the matrix generation will be performed during the initial tracing of the function, assuming the sparsity pattern is constant.

209 Compute device from Inputs

Like PyTorch, Φ_{ML} executes operations on the device where the tensors are allocated. This prevents unintentional copies of tensors as users have to explicitly declare transfer operations. This is unlike TensorFlow, where context managers can be used to specify the target device for code blocks.

214 Custom CUDA Operatorions

 $\Phi_{
m ML}$ provides custom CUDA kernels for specific operations that could bottleneck simulations, such as grid sampling for TensorFlow or linear solves. If available, these will be used automatically in place of the fallback Python implementation.



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