



TorchMorph: CUDA-accelerated Morphological Transforms

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Abstract

Morphological transforms are long-standing tools for shape and mask processing, but the de-facto reference implementation in the Python ecosystem, *i.e.* `scipy.ndimage`, is CPU-only, single-array, and therefore unusable inside a GPU training loop without an expensive device-to-host round trip. GPU vision libraries built on PyTorch cover a narrow subset of these operators, typically restricted to two spatial dimensions and flat structuring elements. We present TORCHMORPH, a lightweight PyTorch extension that closes this gap. TORCHMORPH exposes 22 public operators covering binary morphology, greyscale morphology, exact and approximate distance transforms, and entropy-regularised optimal transport, all implemented as fused CUDA kernels that operate directly on `(B, C, Spatial...)` CUDA tensors with up to eight spatial dimensions. The API deliberately mirrors `scipy.ndimage` argument-for-argument, including border modes, structuring-element origins and pre-allocated outputs, so that existing pipelines port with a change of import. We describe the layered architecture and the kernel designs behind each operator family. Against single-threaded CPU references, batched execution reaches up to 1.1×10^3 times the throughput of `scipy.ndimage` on greyscale morphology and up to $350 \times$ on exact Euclidean distance transforms, while the Sinkhorn solver runs up to $42 \times$ faster than POT. Binary and chamfer operators reproduce their SciPy counterparts exactly, and every float-valued operator agrees with the CPU reference to within 1.8×10^{-6} absolute error. TORCHMORPH is released under the MIT licence at <https://intcomp.github.io/tm>.

Keywords: Morphological Transforms, Distance Transform, Optimal Transport, CUDA, PyTorch, Open Source

1 Introduction

Mathematical morphology supplies some of the oldest and most widely used primitives in image analysis [1, 2]. Erosion, dilation and their compositions underpin denoising, shape decomposition and connected-component post-processing, while the distance transform, an erosion by a parabolic structuring function, underpins skeletonisation and watershed seeding [3, 4], boundary-aware losses [5] and Hausdorff-distance surrogates [6]. In the Python ecosystem these operators are canonically provided by `scipy.ndimage` [7], whose semantics for border modes, structuring-element origins and connectivity conventions have become the de-facto standard that downstream libraries are expected to reproduce.

That reference implementation was designed for a world in which images live in host memory as NumPy arrays [8]. In modern AI-native imaging pipelines, images are represented differently in three specific ways. First, data live on the GPU as PyTorch tensors [9], so every call into `scipy.ndimage` forces a device-to-host copy, a single-threaded CPU computation, and a copy back. Second, data arrive in *batches*: a training step processes tens of volumes at once, and a per-image Python loop over a CPU routine serialises what is naturally an embarrassingly parallel workload. Third, the interesting regime is often three- or four-dimensional (volumetric time series,

multi-channel tomography), where the CPU cost of a morphological sweep grows with the product of all spatial extents.

The obvious response is to use a GPU vision library, but that runs into a coverage problem. Kornia [10] provides differentiable 2-D morphology, but not N -dimensional operators, the full `scipy.ndimage` border-mode matrix, or an exact Euclidean distance transform; cuCIM [11] offers GPU morphology through CuPy, an interoperability layer rather than native `torch.Tensor` operators; MONAI [12] delegates several morphological post-processing steps back to SciPy or CuPy; and OpenCV [13] and scikit-image [14] are host-side and two-dimensional in their fast paths. Entropic optimal transport, increasingly used as a shape- and histogram-comparison loss [15, 16], lives in yet another stack [17, 18]. A practitioner who wants batched GPU morphology, an exact distance transform and a differentiable transport distance must therefore assemble three libraries with three tensor conventions. Table 1 summarises the resulting coverage gap.

TORCHMORPH closes that gap in one place. It brings these operators into the PyTorch tensor world as native, batch-parallel CUDA kernels: 22 public operators spanning binary and greyscale morphology, exact and approximate distance transforms, and entropy-regularised optimal transport. Every one accepts `(B, C, Spatial...)` CUDA tensors of spatial rank up to eight, and mirrors

	GPU	Torch	Batch	N-D	Sem.	EDT	OT
<code>scipy.ndimage</code> [7]	✗	✗	✗	✓	✓	✓	✗
scikit-image [14]	✗	✗	✗	✓	~	✓	✗
OpenCV [13]	~	✗	✗	✗	✗	✓	✗
Kornia [10]	✓	✓	✓	✗	✗	~	✗
cuCIM [11]	✓	✗	✗	✗	~	✓	✗
MONAI [12]	✓	✓	~	✗	✗	~	✗
POT [17]	✓	✓	✓	✗	✗	✗	✓
TORCHMORPH	✓	✓	✓	✓	✓	✓	✓

Table 1: Coverage of the capabilities TORCHMORPH targets. ✓ = supported, ~ = partial or indirect, ✗ = not supported. *Torch* = operates on `torch.Tensor` natively; *Batch* = a single call over a leading batch dimension rather than a Python loop; *N-D* = more than three spatial dimensions; *Sem.* = matches `scipy.ndimage` border modes, structuring-element origins and iteration conventions; *EDT* = exact Euclidean distance transform.

its `scipy.ndimage` counterpart in name, argument order, defaults and boundary behaviour, so porting is a change of import rather than a rewrite, and 3-D and 4-D data are first-class rather than special cases. The operators are fused rather than assembled from generic tensor primitives: one launch per call, with structuring-element geometry resolved on the host, an interior fast path that removes per-axis boundary tests, and CUDA-graph replay for the transport iterations. The library depends only on PyTorch and a CUDA toolchain, is validated element-wise against SciPy and POT, and is released under the MIT licence. Section 2 describes the transforms it covers and the design behind them; Section 3 reports its numerical agreement and throughput against the CPU reference.

2 TorchMorph

2.1 Covered transforms

Figure 1 organises the 22 exported operators into four families plus a structuring-element utility group. Only the bold entries are backed by a dedicated CUDA kernel; the rest are host-side compositions of those primitives, which is why adding a new top-hat variant costs no device code.

Binary and greyscale morphology. Erosion and dilation, and the openings, closings, gradients, Laplacians and top-hats built from them [1, 2, 19], remain the standard tools for cleaning up masks, decomposing shapes and extracting structure at a chosen scale. In deep-learning pipelines they appear as post-processing on predicted segmentations and as label preparation upstream of training [12, 20]. Both run per-step, on batched tensors that already live on the GPU, which is exactly the regime a CPU reference serves badly. TORCHMORPH implements erosion and dilation as fused kernels and derives the remaining eleven operators from them.

Distance transforms. The distance transform labels every foreground element with its distance to the nearest background element. It underpins skeletonisation, watershed seeding and shape descriptors [3, 4], and in learning pipelines it is the ingredient behind boundary-aware and Hausdorff-style losses [5, 6], which recompute a distance field every training step and so put it on the critical path. TORCHMORPH provides the exact Euclidean transform via the separable lower-envelope algorithm [21, 22], chamfer transforms under the chessboard and taxicab metrics [4], and a brute-force transform used as an exactness oracle.

Entropy-regularised optimal transport. Optimal transport compares two distributions by the cheapest way of morphing one into the other, and its entropy-regularised form is solved by the Sinkhorn iteration [23, 15, 16]. It has become a standard geometry-aware loss for histograms, point clouds and attention-style matching problems, where a bin-wise divergence would ignore how far apart the bins are. TORCHMORPH provides a batched Sinkhorn solver in both the scaling and log domains, differentiable with respect to both marginals, so it drops into a training loop as a loss.

2.2 Architecture

2.2.1 API surface

Every operator that consumes an image accepts a CUDA tensor shaped `(B,C,Spatial...)` with $1 \leq \text{spatial rank} \leq 8$, validated by a single shared routine, and takes the same keyword arguments as its SciPy counterpart: `size`, `footprint`, `structure`, `mode`, `cval` and `origin` for greyscale operators; `structure`, `iterations`, `mask` and `border_value` for binary ones; `sampling`, `metric` and the `return_distances/return_indices` pair for distance transforms, with pre-allocated output buffers supported throughout. Iteration semantics match too: `iterations < 1` means *iterate until the result stops changing*, which is how `binary_propagation` and `binary_fill_holes` are built.

```
# before: CPU, one volume at a time
import scipy.ndimage as ndi
out = [ndi.grey_opening(v, size=3) for v in vols]

# after: GPU, whole batch, one launch
import torchmorph as tm
out = tm.grey_opening(vols_cuda, size=3)
```

Listing 1: Porting a SciPy pipeline.

2.2.2 Three layers

The library is organised as in Figure 2. The *Python layer* normalises and validates arguments: it resolves the structuring element from the `structure > footprint > size` priority chain, expands `origin` to a per-axis tuple and range-checks it against the element extent, maps border-mode strings to integer codes, and composes the derived

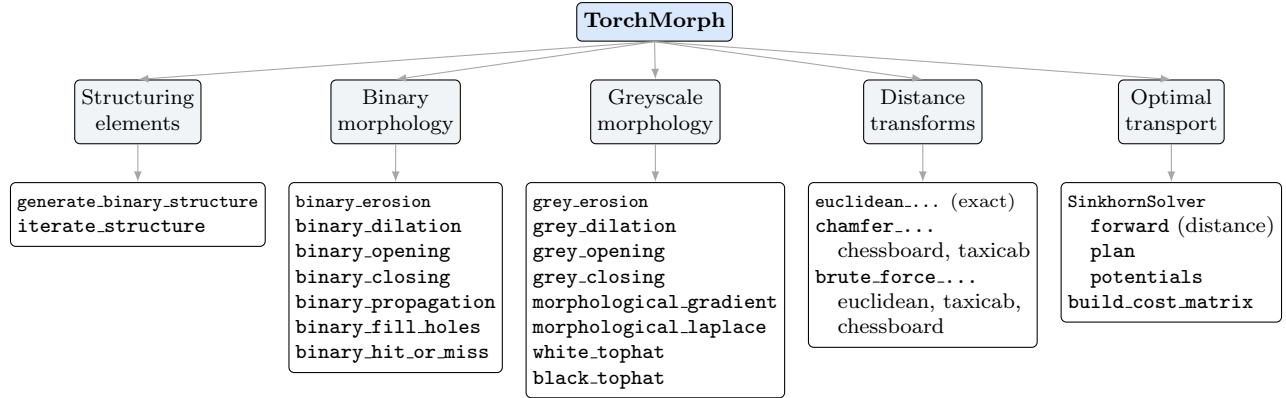


Figure 1: Operator taxonomy of TORCHMORPH. Bold entries are backed by a dedicated CUDA kernel; the remainder are host-side compositions of those primitives. All image operators accept (B, C, Spatial...) CUDA tensors with up to eight spatial dimensions.

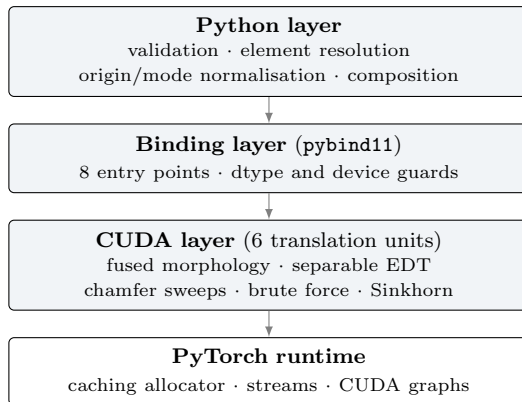


Figure 2: The three implementation layers. SciPy-compatible argument conventions are confined to the Python layer; the CUDA layer sees only normalised geometry.

operators from the two primitives. This layer is where SciPy compatibility is defined, and deliberately the only place that knows about argument conventions. The *binding layer* is a single `pybind11` module exposing eight kernel entry points compiled from six CUDA translation units; keeping that surface small means derived operators cost no extra device code. The *kernel layer* holds the CUDA implementations, with two decisions recurring across it: all geometry that can be resolved on the host is resolved on the host, and every kernel is written against a runtime spatial rank with a compile-time bound, so co-ordinate scratch space stays in registers; the Python layer caps that rank at eight.

2.3 Kernel design

2.3.1 Fused morphology kernel

A naive GPU morphology kernel recomputes, for every output element and every structuring-element position, a full N -dimensional coordinate mapping with a per-axis boundary test. For a 3^3 element in 3-D that is 27 mapped

coordinates per voxel, each requiring the linear index to be decomposed along every axis before it can be bounds-checked.

TORCHMORPH avoids this in two ways. On the host, the structuring element is flattened once into a list of active entries; for each we store the per-axis offset *and* the precomputed flat offset against the input’s spatial strides, and inactive footprint positions never reach the device.

On the device, each thread first tests whether its output coordinate is *interior*, using per-axis offset extrema computed on the host. Interior threads, the overwhelming majority, take a fast path that adds the precomputed flat offset directly to the linear index, with no per-axis arithmetic and no boundary test at all. Only threads within one element radius of a face take the general path, which resolves out-of-bounds neighbours as its SciPy counterpart does: the greyscale kernel implements all five border modes (**constant**, **reflect**, **nearest**, **mirror**, **wrap**), the binary kernel a single **border_value**.

Erosion and dilation are instantiations of one templated kernel parameterised by a functor supplying the identity element and the combining rule, and the binary kernel adds a **done** predicate so a thread can break out of the reduction as soon as the result is decided, false for erosion and true for dilation, which is a substantial saving on sparse masks.

2.3.2 Distance-transform kernels

The EDT uses the separable lower-envelope formulation, in which each one-dimensional pass computes the lower envelope of a family of parabolas in a single sweep [21]. That sweep is inherently sequential, so TORCHMORPH assigns one thread block per scanline: a single thread builds the envelope in shared memory while the whole block co-operates on loading the line and, afterwards, on the query phase, where each output position binary-searches the intersection array in parallel. Parallelism comes from the number of scanlines, which is ample: for B batch items of

C channels and extent n^d it is BCn^{d-1} per pass.

Three paths exist: a 2-D specialisation for extents up to 2048 that keeps the row pass fully contiguous and fuses the final square root into the column pass; a general path that transposes the active axis to be innermost so every pass sees contiguous memory; and a fallback that spills the envelope stack to a lazily allocated global buffer when a scanline exceeds the shared-memory budget. Nearest-background indices are propagated through the passes on request, and an image with no background is handled by the same virtual out-of-bounds convention SciPy uses.

The chamfer transform is implemented as dimension-separable forward and backward sweeps, with extra diagonal passes for the chessboard metric. The brute-force transform stages background coordinates through shared memory in tiles of 256 and is templated over both metric and spatial rank, so the coordinate loop is fully unrolled and metric selection costs no branch; its quadratic cost confines it to validating the separable transforms and to the one case they miss: the chamfer entry point takes no `sampling` argument, so anisotropic chessboard and taxi-cab distances are reachable only this way.

2.3.3 Batch-tiled Sinkhorn solver

The transport module solves the Sinkhorn iteration for a batch of n histogram pairs of dimension d sharing one $d \times d$ cost matrix, and three characteristics of that workload drive the design.

The matrix is shared across the batch: rather than the generic batched matrix–vector product, which reads the d^2 matrix once per batch item, TORCHMORPH assigns one block per (row, batch-tile) pair with a tile of eight items, streams the matrix row once and applies it to all eight scaling vectors held in registers, dropping matrix traffic by the tile factor.

The log-domain update needs only one pass: where the textbook log-sum-exp makes two passes over the row, one for the maximum and one for the shifted sum, TORCHMORPH maintains a running maximum and a rescaled running sum in a single pass and reduces the resulting pairs across the block with a merge operator that is well defined on the empty state, so an all-zero marginal pins the potential to $-\infty$ instead of producing NaN.

Long runs are launch-latency bound: each iteration is two small kernels, so from 100 requested iterations upwards TORCHMORPH captures a chunk of 25 into a CUDA graph [24] and replays it. Because the iterations are fixed-point steps on ping-pong buffers, executing extra iterations is always safe, and the graph path degrades gracefully to plain launches when capture is unavailable.

Gradients come from a custom autograd function returning the centered dual potentials, which by the envelope theorem are the exact gradients of the entropic transport cost with respect to the marginals [16]; the potentials are centred and stashed during the forward pass, so the

backward pass is a single broadcast multiply and never differentiates through the iteration.

2.4 Testing and reference alignment

Matching `scipy.ndimage` is a claim about behaviour, so it is checked by differential testing rather than asserted. The suite is 78 test functions across five modules.

Oracle agreement accounts for most of it: every exported operator is compared elementwise against its reference (`scipy.ndimage` for morphology and distance transforms, POT for transport) over 2-D, 3-D and higher-rank inputs, batch and channel combinations, non-contiguous and transposed layouts, anisotropic sampling, every border mode, and asymmetric structuring elements with shifted origins. The reference is applied sample by sample to the same array the GPU receives, so batching cannot mask a per-item discrepancy. Where no reference settles the question, *invariants* take over: transport plans must reproduce both marginals, returned indices must point at a genuine nearest background element, and the analytic gradient is checked against a finite-difference directional derivative.

The remainder guards the seams. *Cross-implementation* tests solve one problem along every path the library can take: fused kernels against the pure-torch fallback, float32 against float64, CPU against GPU, CUDA-graph replay against plain launches, early-stopped against fully iterated. The results must coincide. *Contract* tests, a quarter of the suite, assert failure on spatial rank above eight, mismatched origin, mask or output shapes, unknown modes and metrics, and non-CUDA input. *Runtime* tests issue work on a side stream and a non-default device, checking that kernels honour the caller’s stream and device rather than the defaults.

3 Results

Protocol. All comparisons use `scipy.ndimage` as the reference, except for transport, which uses POT [17]. All measurements come from a single machine. *Hardware:* NVIDIA GeForce RTX 4090 D GPU (Ada Lovelace, 48 GB, driver 580.173.02), $2 \times$ Intel Xeon Gold 6330 CPU (56 cores / 112 threads, 2.00 GHz), and 128 GB of system memory. *Software:* Ubuntu 24.04.3 LTS, CUDA 12.4, Python 3.12.13, PyTorch 2.6.0+cu124, NumPy 2.5.1, SciPy 1.18.0, and POT 0.9.6.post1. TORCHMORPH extensions were compiled with `-O3` and `--use_fast_math`. The SciPy and POT baselines are single-threaded and therefore occupy one CPU core, while every GPU result uses a single device. Accuracy is measured by transferring the GPU result to the host and comparing against the reference output on the identical input; we report the maximum absolute error $\max |y - \hat{y}|$ and the relative error $\|y - \hat{y}\|_2 / \|y\|_2$. Timings are collected with

Operator family	Reference	Max abs. err.	Rel. ℓ_2 err.
Binary morphology	<code>ndimage</code>	0	0
Grey morphology	<code>ndimage</code>	4.77×10^{-7}	2.24×10^{-8}
Euclidean DT	<code>ndimage</code>	2.06×10^{-7}	6.77×10^{-9}
Chamfer DT (chessb.)	<code>ndimage</code>	0	0
Chamfer DT (taxicab)	<code>ndimage</code>	0	0
Brute-force DT (ℓ_2)	<code>ndimage</code>	2.06×10^{-7}	6.71×10^{-9}
Sinkhorn distance	POT	1.75×10^{-6}	8.82×10^{-8}

Table 2: Worst-case numerical error over all tested configurations against the CPU reference implementations. Binary morphology and chamfer distance transforms match `scipy.ndimage` exactly in the tested cases, while float-valued operators differ only at float32-level numerical precision. Sinkhorn distance is compared against POT.

`torch.utils.benchmark`, taking the median of blocked auto-ranged measurements with a one-second minimum run time and per-item normalisation, after warm-up and with an explicit `torch.cuda.synchronize()` inside the timed region. Because the point of the library is batching, we report two GPU columns: $1\times$, a Python loop calling the operator once per batch item, the honest analogue of a SciPy loop, and *batch*, a single call on the whole batch. Speed-up is SciPy time divided by batched GPU time per item. Every table is reproduced by the scripts in [benchmark/](#).

Numerical agreement. Table 2 shows close agreement with the CPU references. Binary morphology and chamfer distance transforms match `scipy.ndimage` exactly; all float-valued operators have worst-case absolute and relative ℓ_2 errors below 1.8×10^{-6} and 9×10^{-8} , respectively. Small discrepancies arise from float32 GPU arithmetic and `--use_fast_math`. NaN propagation is the only documented behavioral difference from SciPy. Correctness is covered by 78 test functions, parametrised over operator, spatial rank, input shape, origin, sampling, requested outputs, device and solver variant, with `scipy.ndimage` as the oracle for morphology and distance transforms and POT for transport. The suite runs on every push and pull request through a self-hosted continuous-integration runner equipped with a physical CUDA device, so each change is checked against the CPU references by executing the actual kernels rather than a mocked or CPU-only code path; a second hosted runner enforces formatting and static checks.

Throughput. Table 3 reports per-input throughput for morphology and distance transforms. The $B = 1$ results capture single-input GPU execution, where launch overhead can dominate, while larger batches expose the parallel regime targeted by TORCHMORPH. The batching benefit generally decreases as individual inputs become large enough to better utilize the GPU.

Table 3: Per-input throughput across representative operators, input sizes, and batch sizes. Throughput is reported in inputs/ms (higher is better), where one input denotes one 2-D image or one 3-D volume. TM = TORCHMORPH; its values are rounded to one decimal place, while the SciPy column keeps three, since one decimal would collapse most of its entries to 0.0.

Operator	Size	Throughput (inputs/ms) $\uparrow$				
		SciPy	TM			
			$B=1$	$B=2$	$B=4$	$B=8$
Binary erosion	256^2	0.865	6.9	13.7	27.8	55.6
	1024^2	0.056	6.8	12.8	22.2	35.7
Binary dilation	256^2	0.851	6.7	13.5	27.0	52.6
	1024^2	0.057	6.7	12.5	21.7	35.7
Grey erosion	256^2	0.740	16.1	32.3	62.5	83.3
	1024^2	0.040	13.7	23.3	34.5	45.5
Grey dilation	256^2	0.756	10.3	30.3	58.8	111.1
	1024^2	0.040	13.2	22.2	33.3	45.5
EDT 2-D	256^2	0.160	13.3	25.6	30.3	40.0
	1024^2	0.011	2.1	2.7	2.8	2.9
EDT 3-D	64^3	0.031	6.3	8.3	9.3	10.8
	128^3	0.003	1.4	1.4	1.5	1.5
CDT chessboard	256^2	0.585	4.7	9.4	18.2	33.3
	1024^2	0.037	1.5	2.6	4.3	5.9
CDT taxicab	256^2	0.637	5.5	10.9	20.0	43.5
	1024^2	0.041	1.7	3.4	6.6	11.6
BFDt (ℓ_2)	256^2	< 0.001	1.1	1.2	1.2	1.2

Morphology and distance transforms. Table 3 reports per-input throughput for representative morphological operators and distance transforms. Across most operators, throughput increases substantially with batch size, reflecting the fact that a single input does not supply enough parallel work to saturate the GPU. Grey dilation on 256^2 images, for instance, improves from 10.31 inputs/ms at $B = 1$ to 111.11 inputs/ms at $B = 8$ (a $10.8\times$ gain), while grey erosion improves from 16.13 to 83.33 inputs/ms ($5.2\times$). Binary morphology follows the same pattern, reaching 36–56 inputs/ms at $B = 8$ depending on the operator and image size.

The benefit of batching diminishes markedly for computationally heavier transforms, where a single large input already supplies enough parallelism to approach device saturation on its own. EDT on 1024^2 images improves only modestly, from 2.11 to 2.86 inputs/ms ($1.4\times$), and EDT on 128^3 volumes remains essentially flat at approximately 1.4 inputs/ms regardless of batch size. CDT retains a clear batching benefit, particularly for smaller inputs, whereas BFDt shows only a marginal gain, from 1.15 to 1.25 inputs/ms. SciPy’s throughput on BFDt is extremely low because it relies on a brute-force reference implementation, so this comparison should be read as a correctness oracle rather than as a benchmark against an optimized CPU distance-transform implementation.

Table 4: Entropic optimal transport against POT on 2-D grid problems with $d = 32^2$ bins (TM = TORCHMORPH). The 1000-iteration row uses CUDA-graph replay. Timings are rounded to one decimal place; speed-ups are computed from the unrounded values.

Configuration	POT (ms)	TM (ms)	Speed-up	Rel. err. (plan)
scaling, 100 it.	23.0	1.2	18.8×	9.21×10^{-4}
scaling, 1000 it.	229.7	10.1	22.8×	4.45×10^{-7}
log-domain, 200 it.	35.4	3.0	11.8×	4.55×10^{-5}
batch $n = 16$, 100 it.	367.3	8.7	42.4×	1.54×10^{-3}

Optimal transport. Table 4 compares the Sinkhorn solver against POT [17] on grid-structured problems, where the cost matrix is the pairwise ℓ_2 distance between the points of a 32×32 grid. We also report the relative error of the recovered transport plan. Three configurations are of particular interest: the scaling form at moderate regularisation, the log-domain form at small regularisation where the scaling form underflows, and a batched run that exercises the tiling described in Section 2.3.3.

Threats to the comparison. The speed-ups therefore compare a GPU against one CPU core, not a well-parallelised CPU implementation; we report per-item times and the $B = 1$ column so the batching effect can be separated from the device effect. The brute-force timings are the cost of a correctness oracle, not a performance claim.

4 Conclusion

TORCHMORPH provides batch-parallel, N -dimensional CUDA implementations of binary and greyscale morphology, exact and approximate distance transforms, and entropy-regularised optimal transport, behind an API that mirrors `scipy.ndimage`. Its contribution is availability rather than algorithmic novelty: classical algorithms implemented once, correctly, against the tensor conventions modern imaging pipelines actually use, and validated against the references the community already trusts. That turns these operators from an offline preprocessing step into something that can sit inside a training loop [5, 6].

Three limitations remain. The morphology and distance kernels are forward-only; only the transport module is differentiable, though erosion and dilation admit subgradients routed to the arg-min/arg-max position. They also require CUDA, the intended fallback being SciPy itself, whereas the transport module drops back to pure torch on CPU. And they compute in float32 under `--use_fast_math`, with NaN propagation not guaranteed to match the reference. Future work is autograd for the morphological operators and a wider set of connected-component and reconstruction operators.

Acknowledgements

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