



DeepEverest: Accelerating Declarative Top-K Queries for Deep Neural Network Interpretation

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ABSTRACT

We design, implement, and evaluate DeepEverest, a system for the efficient execution of *interpretation by example* queries over the activation values of a deep neural network. DeepEverest consists of an efficient indexing technique and a query execution algorithm with various optimizations. We prove that the proposed query execution algorithm is instance optimal. Experiments with our prototype show that DeepEverest, using less than 20% of the storage of full materialization, significantly accelerates individual queries by up to 63× and consistently outperforms other methods on multi-query workloads that simulate DNN interpretation processes.

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The source code, data, and/or other artifacts have been made available at <https://github.com/uwdb/deepeverest>.

1 INTRODUCTION

Deep neural networks (DNNs) are increasingly used by machine learning (ML) applications. When training and deploying DNNs, interpretation is important for researchers to understand what their models learn. DNN interpretation is a relatively new field of research, and techniques are evolving. While many new approaches are developed, they often do not scale with the size of the datasets and models [47]. The problem we address in this paper is the efficient execution of a common class of DNN interpretation queries.

The fundamental building blocks of DNN interpretation are neurons. Each neuron outputs an activation value as the input is propagated through the network. DNN interpretation techniques often perform analysis on these activation values of neurons [5, 6, 13, 41, 61, 63]. When DNNs are trained on tasks such as image classification or scene synthesis, there emerge individual neurons

and groups of neurons that match specific human-interpretable concepts [6, 13, 62], such as “human faces” and “trees”.

To understand what individual neurons and groups of neurons learn and detect, researchers often ask *interpretation by example* queries [29], which are important constituents of the class of post-hoc interpretation methods that are applied to trained models, as opposed to methods that achieve interpretability by restricting model complexity [38]. A widely used *interpretation by example* query is, “find the top-k inputs that produce the highest activation values for an individual neuron or a group of neurons” [12, 14, 22, 34, 51, 58, 59, 62]. Another common query is, “for any input, find the k-nearest neighbors in the dataset using the activation values of a group of neurons based on the proximity in the latent space defined by the group of neurons” [2, 9, 24, 37, 39, 44, 55]. These queries help with investigating and understanding the functionalities of neuron groups by tying those functionalities to the input examples in the dataset. Moreover, these queries are staple techniques for verifying hypotheses of what groups of neurons learn and detect. For instance, Mikolov et al. ask the latter query to find the nearest neighbors of words in the latent space to examine the learned representations after training the word2vec model [37]. As a concrete example, consider a DNN trained to classify images. A user may be interested in understanding what parts of a dog image cause the DNN to predict its class. The user may inspect the maximally activated neurons of different layers in the DNN based on the conjecture that groups of maximally activated neurons act as semantic detectors of features in the image (e.g., floppy ears). To investigate whether these neurons exhibit similar behavior for other images, the user may then ask for the most similar images to the sample image based on the activation values of a group of neurons.

This paper presents a system called DeepEverest that focuses on accelerating the aforementioned two kinds of queries: (1) find top-k inputs that produce the highest activation values for a user-specified group of neurons, and (2) find the top-k most similar inputs based on a given input’s activation values for a user-specified neuron group. A group of neurons consists of one or more neurons within a layer of the DNN. We call the first type of query the *top-k highest* query and the second type of query the *top-k most-similar* query.

Executing these *interpretation by example* queries efficiently with low storage overhead is challenging. A baseline approach is to materialize the activation values for all inputs and all neurons. However, this approach requires significant storage space. For example, storing all the activations uncompressed of ResNet50 for 10,000 images occupies 1.35 TB of disk storage. At the other

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extreme, computing all activation values at query time imposes no storage overhead, but is compute-intensive and extremely slow because it requires DNN inference to compute the activation values on the entire dataset at query time. For instance, answering a *top-k most-similar* query that targets a relatively late layer of ResNet50 on a dataset of 10,000 images takes more than 120 seconds, which renders the DNN interpretation process tedious.

Further, although the target query is a *k*-nearest neighbor (KNN) search, existing approaches that accelerate KNN queries are not applicable. KNN methods rely on building efficient data structures such as trees [7, 32, 42] or hash tables [3, 10] in advance for faster query execution later. One could try to build a single, large, multidimensional data structure for all neurons in each layer. However, such an index would not perform well because of its very large dimensionality. DNNs frequently have layers with multiple thousands of neurons, thus dimensions. One could build data structures for all possible neuron groups that a user could query. However, this would either limit the user to a small set of possible queries or would be prohibitively expensive both in time and storage because the number of possible neuron groups grows exponentially with the number of neurons in each layer. Additionally, in all cases, precomputing and storing all activation values in such data structures would add prohibitive storage overhead.

While many systems have been developed to enable various forms of DNN interpretation [2, 22, 23, 26, 45, 47], none supports flexible and efficient *interpretation by example* queries. In prior work [35], we investigated the use of sampling for model diagnosis. That work, however, focused only on aggregate queries. The closest work to DeepEverest is MISTIQUE [53]. It introduces storage techniques such as compression and quantization, which are orthogonal to DeepEverest and could complement our approach. It is, however, possible to use some of MISTIQUE’s techniques as a caching algorithm, which we compare against in our experiments.

In DeepEverest, we design an index called the Neural Partition Index (NPI), and an efficient query execution algorithm, called the Neural Threshold Algorithm (NTA), which has low storage overhead, reduces the number of activation values that must be computed at query time, and guarantees the correctness of the query results. NTA builds on the classic threshold algorithm (CTA) [11], which supports *top-k* queries that target arbitrary neuron groups. However, CTA requires the computation of the activations of all inputs in the dataset at query time. Because the time to compute the activations by performing DNN inference (not the calculation of the *top-k* result) dominates the end-to-end query time, CTA would not accelerate our target queries. We argue that for any algorithm to improve query time, it must reduce the number of inputs on which DNN inference is run at query time. DeepEverest achieves this reduction of DNN inference at query time while keeping the storage overhead low by building NPI and using it in NTA. Rather than store the raw activations for all neurons, NPI partitions the inputs and stores a small amount of information per-partition that is useful when deciding which activation values to recompute at query time. NTA then uses insights from CTA to decide when to terminate as it incrementally computes activation values using DNN inference only for small subsets of inputs as needed to answer the query. We analyze NTA and show that it is instance optimal for finding the query results of our target queries.

In addition to its fundamental approach, DeepEverest also includes several important optimizations: (1) incremental indexing to avoid large preprocessing overhead; (2) Maximum Activation Index (MAI) to accelerate *top-k most-similar* queries that target maximally activated neurons and *top-k highest* queries; (3) automatic configuration selection; and (4) Inter-Query Acceleration (IQA), which further speeds up sequences of related queries.

In summary, the contributions of this paper are:

- We propose, design, and implement a system called DeepEverest that includes an efficient index structure and an instance optimal query execution algorithm that accelerates *interpretation by example* queries for DNN interpretation while keeping the storage overhead low (Sections 4.2, 4.3, 4.4, 4.5 and 4.6).
- We develop additional optimizations for DeepEverest that further accelerate individual queries (Section 4.7.1), automatically select a good configuration for the system (Section 4.7.2), and accelerate sequences of related queries (Section 4.7.3).
- We evaluate DeepEverest on benchmark datasets and models (Section 5). We demonstrate that DeepEverest, using less than 20% of the storage of full materialization, significantly accelerates individual *interpretation by example* queries by up to 63.5× and consistently outperforms other methods on multi-query workloads that simulate DNN interpretation processes.

2 PRELIMINARIES

A DNN consists of layers composed of units, called neurons, connected by edges with associated weights. Inputs to a DNN are propagated through the layers. The output of a neuron is a linear combination of its inputs and their associated edge weights that is optionally transformed by a nonlinear activation function. The output of each neuron for a given input is called its *activation value* or *activation*. DNN interpretation often involves the study of these activations. Typical questions that researchers may ask include the *interpretation by example* queries mentioned in Section 1. These queries enable researchers to reason about what the DNN learns and identify how groups of neurons match human-interpretable concepts. In this paper, we address the problem of enabling fast queries over activations in a DNN. Conceptually, a DNN and an input dataset can be described by the relations `Neuron(neuronID, layerID, ...)` and `Artifact(inputID, neuronID, activation)`.

DeepEverest supports two fundamental classes of queries over activations: *top-k highest* queries that find the *top-k* inputs that produce the highest activations for a user-specified group of neurons and *top-k most-similar* queries that find the *top-k* inputs that are most similar to a user-specified target input based on the activations of a user-selected group of neurons. The rank of an input is decided by a user-specified distance function (or a default function), `DIST`. Based on the user-selected neuron group, for *top-k highest* queries, `DIST` takes as input a set of activations and measures their magnitude. For *top-k most-similar* queries, `DIST` measures the distance between the input and the target input, and it takes as input a set of absolute differences between the input’s activations and the target input’s activations. Note that *top-k highest* queries can be considered as *top-k most-similar* queries with a hypothetical target input whose activations are infinite for all neurons. This distance function `DIST` must be monotonic, i.e., $\text{DIST}(x_1, x_2, \dots, x_n) \leq \text{DIST}(x'_1, x'_2, \dots, x'_n)$ whenever $x_i \leq x'_i$ for each i . Monotonicity

is satisfied by common distance functions, such as l_p -distances, cosine distance (once transformed to normalized l_2 -distance), and weighted distances like Mahalanobis distance, among others. The default in DeepEverest is l_2 -distance.

3 RELATED WORK

DNN Interpretation. Many approaches have been proposed to interpret the internals of DNNs [2, 5, 6, 9, 12–14, 22, 24, 34, 39, 41, 44, 51, 55, 58, 59, 61–63]. These approaches ask *interpretation by example* queries that return the most similar inputs with respect to the activations of a group of neurons of a given input, or inputs that maximally activate a group of neurons. They motivate the design of DeepEverest, which does not invent new interpretation methods but rather builds novel indexes and algorithms that accelerate the execution of these commonly asked queries.

Systems for Machine Learning. Many systems have been proposed to support efficient ML [36, 49, 54, 57]. DeepEverest falls into the group of systems that support efficient model diagnosis and interpretation. A number of systems like ModelTracker [2], CNNVis [30], and others [8, 22, 23, 25, 28, 33, 58] support visual inspection of ML models and features. These systems could utilize DeepEverest to accelerate some of the queries used to build the visualizations. DeepBase [47] lets users identify neurons that have statistical dependencies with user-specified hypotheses. However, it does not support *interpretation by example* queries. MISTIQUE [53] accelerates the examination of neuron activations by using storage techniques such as quantization to reduce the storage overhead while sacrificing query accuracy. These techniques are orthogonal to DeepEverest’s and could be incorporated in DeepEverest to further reduce the storage overhead. MISTIQUE also proposes a cost model that captures the trade-off between materialization and recomputation of the activations for different layers and makes materialization decisions accordingly. None of these systems have addressed the problem of accelerating *interpretation by example* queries well because none of them reduce the number of activation values computed during query execution as DeepEverest does.

Nearest Neighbor Search. Our target query is a k -nearest neighbor (KNN) search. While many methods exist for exact [7, 15, 32, 42] and approximate [3, 10, 20, 21, 56] KNN, the challenge in this paper is different. These KNN methods must know what dimensions will be queried before constructing data structures in that space. In our problem, the dimensions of the KNN search are defined by the neuron group specified only at query time.

Top-K Query Processing. Top- k query processing is formalized by the seminal work on the threshold algorithm [11]. The algorithm scans multiple sorted lists and maintains an upper bound for the aggregate score of unseen objects. Each newly seen object is accessed (by random access) in every other list, and the aggregate score is computed by applying the scoring function to the object’s value in every list. The algorithm terminates after k objects are seen with scores greater than or equal to the upper bound. Many follow-up approaches propose approximation, optimizations, and extensions [1, 4, 16, 19, 43, 52, 60], but they assume that accesses are available to the underlying data sources. This assumption does not hold in our problem. Storing (on disk) or computing (at query time) the activations required for the top- k queries incurs prohibitive storage overhead or computation overhead respectively.

4 DEEVEREST

In this section, we first consider baseline approaches, then describe how DeepEverest improves upon these baselines to accelerate query execution while keeping the storage overhead low.

4.1 Baselines

We first discuss baseline approaches and explain why applying CTA or any KNN algorithm would not improve the query time.

PreprocessAll. The first baseline, *PreprocessAll*, has a high storage cost. It performs DNN inference on the entire dataset and stores all the activations for all neurons ahead of time. It executes queries by loading the previously stored activations of the neuron group for all inputs from disk and maintaining a top- k result set.

ReprocessAll. The second baseline, *ReprocessAll*, has a high computation cost. It has no storage overhead and performs no preprocessing. It executes queries by computing the activations of the layer being queried by DNN inference on all inputs and maintaining a top- k result set as it performs the computation.

LRU Cache. The third baseline, *LRU Cache*, is a disk cache that has a fixed storage budget with a least-recently-used (LRU) replacement policy. This strategy strikes a balance between the storage overhead of *PreprocessAll* and the computation overhead of *ReprocessAll*. *LRU Cache* maintains a fixed-sized disk cache that stores the activations for queried layers. A query is executed as in *PreprocessAll* if the activations of the queried layer are present in the disk cache. Otherwise, it is executed as in *ReprocessAll*. After that, the activations of the queried layer are persisted to the disk cache. When the size of the disk cache exceeds the storage budget, the cache evicts the activations of the least recently used layer.

Priority Cache. The final baseline, *Priority Cache*, is a technique adapted from MISTIQUE [53]. It has a fixed-sized disk cache to store the activations for some layers. As a preprocessing step, it uses the storage cost model from [53] to pick which layers to store, assuming that each layer will be queried the same number of times. Under the storage budget, this cost model prioritizes the layers that save the most query time per GB of data stored. It performs DNN inference on every input and stores the activations for the layers selected ahead of time. A query is executed as in *PreprocessAll* if the activations of the queried layer are present in the disk cache. Otherwise, the query is executed as in *ReprocessAll*.

CTA could be applied to each baseline by first using the materialized or recomputed activations to construct the *Artifact* table defined in Section 2. *Artifact* is then used to construct a relation in which each row represents an input, and each column represents a neuron and contains the absolute difference between the activation of that row’s input and the target input’s activation on the column’s neuron. CTA can run after sorting the absolute differences in each column in ascending order, using any monotonic norm of the absolute differences as the aggregation function.

However, applying CTA (or any KNN algorithm) on top of each of the baselines would not improve the query time. Using it along with *ReprocessAll* would not help because *ReprocessAll* requires running DNN inference on the entire dataset to compute *Artifact* at query time. Similarly, applying it on *PreprocessAll* would not improve the query time because generating the relation of absolute differences requires a full scan over *Artifact* before CTA can be

Table 1: Query time breakdown for baselines for a $top-k$ most-similar query on *ImageNet-ResNet50* (query: *SimHigh*, neuron group size: 3, layer: *late*; detail in Section 5.1).

Method	<i>ReprocessAll</i>	CTA [11]	<i>K-D Tree</i> [7]	<i>Ball Tree</i> [42]
Total query time	121.6 s	121.6 s	121.8 s	121.7 s
DNN inference time	121.4 s	121.4 s	121.4 s	121.4 s

applied. The $top-k$ result could already be computed during the full scan. Further, applying it on *PreprocessAll* incurs this strategy’s prohibitive storage overhead. Applying it on top of *LRU Cache* and *Priority Cache* would not improve the query time for the same reasons as *PreprocessAll* (for layers in the cache) and *ReprocessAll* (for layers not in the cache). Given the prohibitive storage overhead of *PreprocessAll*, any existing method that supports queries for arbitrary neuron groups must compute the activations of the neuron group for all inputs at query time. Table 1 shows the query time breakdown for various baselines. The total query time consists of the time for DNN inference to compute the activations of the queried neuron group, the time for building the data structure required for each method (it cannot be computed ahead of time because the neuron group for a query can be arbitrary), and the time to obtain the $top-k$ result. As the results show, DNN inference is the bottleneck of query execution. Therefore, any method that does not reduce the number of inputs fed into the DNN at query time will perform similarly to *ReprocessAll*. Hence, the query time of *ReprocessAll* can represent that of these more advanced methods.

4.2 Overview of DeepEverest

As described above, directly applying CTA does not improve the query time because Artifact must be fully computed for the neuron group at query time, which requires DNN inference on all inputs. Query execution can be significantly accelerated by avoiding running the DNN on inputs that will not be one of the $top-k$ results.

We design and build a novel index, called the Neural Partition Index (NPI) (Section 4.3), and a query execution algorithm, called the Neural Threshold Algorithm (NTA) (Section 4.4). NTA is a modified threshold algorithm. In contrast to CTA, NTA does not require all activations of all inputs before it starts. It utilizes NPI to progressively access and perform DNN inference on only the inputs that are possibly in the $top-k$ results. NTA overcomes the bottleneck of query execution, DNN inference, by reducing the number of inputs on which DNN inference is performed at query time, while guaranteeing the correctness of the $top-k$ results and introducing only tolerable storage overhead. Moreover, we show the instance optimality of NTA (Section 4.5) and propose various additional optimizations (Sections 4.6 and 4.7).

4.3 Neural Partition Index (NPI)

A key goal of DeepEverest is to support queries that target arbitrary neuron groups. Existing partitioning or indexing methods (e.g., *K-D Tree*, *locality-sensitive hashing*) must know which neuron group will be queried ahead of time and construct data structures in that space. In contrast, DeepEverest constructs indexes for each neuron separately and therefore is able to answer queries for arbitrary neuron groups. The activations in a DNN can conceptually be

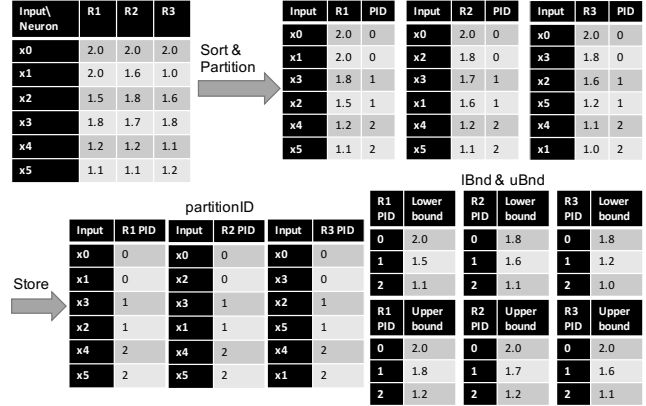


Figure 1: An example of building the Neural Partition Index of three neurons, $R1, R2, R3$, for six inputs, $x0, \dots, x5$.

represented by the Artifact relation introduced in Section 2. Recall that `neuronID` is the identifier for a neuron in the DNN and `inputID` is the identifier for an input in the dataset. DeepEverest conceptually builds an index on the search key (`neuronID`, `activation`) and supports queries that return the `inputIDs` for a given `neuronID` and range of activation values. To avoid materializing activations, DeepEverest builds an index on (`neuronID`, `PID`) instead, where `PID` (partitionID) is the identifier of a range-partition over activation values for a neuron. DeepEverest builds equi-depth partitions instead of equi-width partitions because the activation values are usually highly skewed, and equi-depth partitions adapt better to skewed distributions. Partition 0 contains the largest activations. The index then supports efficient lookups for a given (`neuronID`, `PID`) combination. The index returns the set of `inputIDs` whose activations for the given `neuronID` belong to the partition identified with `PID`. Moreover, the index also supports queries that return the `PID` for a given (`neuronID`, `inputID`) combination. We call this structure the Neural Partition Index (NPI) and denote the queries with `getInputIDs(neuronID, PID)` and `getPID(neuronID, inputID)`. Additionally, in NPI, for each partition, DeepEverest stores the *lower bound* and *upper bound* of the activations in that partition and supports queries that ask for them. We denote such queries with `lBnd(neuronID, PID)` and `uBnd(neuronID, PID)`. The number of partitions, $nPartitions$, is configurable and discussed further in 4.7.2.

There are two approaches to implementing the index. The first approach would be to maintain a set of buckets, each identified with a unique (`neuronID`, `PID`) combination as the key, and, for each bucket, maintain a list of `inputIDs`. The second approach, which DeepEverest uses, is to maintain a list of (`neuronID`, `inputID`) pairs as keys, and, for each entry, store the `PID`. `neuronID`, `inputID`, and `PID` are integers, so rather than building a B-tree or a hash index over the keys, we create an optimized index structure using an array where `neuronID` and `inputID` act as offsets for lookups in the array. This enables DeepEverest to only store the values and therefore avoid the cost of storing the keys. Figure 1 illustrates NPI for an example dataset with three partitions.

During preprocessing, DeepEverest runs DNN inference on all inputs once to build NPI for every neuron, which requires sorting. The time complexity to compute NPI once the activations are

dPar				ord			
PID\Neuron	R1	R2	R3	c\Neuron	R1	R2	R3
0	0.9	0.7	0.6	0	2	2	1
1	0.4	0.5	0.0	1	1	1	2
2	0.0	0.0	0.1	2	0	0	0

Figure 2: *dPar* and *ord* for the execution of NTA for the example query that finds the most similar inputs to x_5 based on the activations of $\{R1, R2, R3\}$. For neuron i , c is the index of the PIDs in *ord*(i), e.g., when $c=0$, $ord(R1, c)=2$.

toRun ($c = 0$)				toRun ($c = 1$)			
Neuron	R1	R2	R3	Neuron	R1	R2	R3
toRun	{x4, x5}	{x4, x5}	{x2, x5}	toRun	{x2, x3}	{x1, x3}	{x1, x4}

(a) Build *toRun*

dist(s, x, G) ($c = 0$)			dist(s, x, G) ($c = 1$)		
Input newly computed	x2	x4	Input newly computed	x1	x3
dist	1.5	0.3	dist	1.6	1.9

top ($c = 0$)			top ($c = 1$)		
$(x, dist(s, x, G))$	{x4, 0.3}	{x2, 1.5}	$(x, dist(s, x, G))$	{x4, 0.3}	{x2, 1.5}

(b) Perform DNN inference to compute *dist* and update *top*

minBoundary, maxBoundary ($c = 0$)				minBoundary, maxBoundary ($c = 1$)			
Neuron	R1	R2	R3	Neuron	R1	R2	R3
(min, max)	(1.1, 1.2)	(1.1, 1.2)	(1.2, 1.6)	(min, max)	(1.1, 1.8)	(1.1, 1.7)	(1.0, 1.6)

F, V ($c = 0$)				F, V ($c = 1$)			
Neuron	R1	R2	R3	Neuron	R1	R2	R3
(F, V)	($\infty, 1$)	($\infty, 1$)	(1, 1)	(F, V)	($\infty, 1$)	($\infty, 1$)	($\infty, 1$)

minDist ($c = 0$)				minDist ($c = 1$)			
Neuron	R1	R2	R3	Neuron	R1	R2	R3
minDist	0.1	0.1	0.0	minDist	0.7	0.6	0.4

$1.5 > t = 0.1 + 0.1 + 0.0 = 0.2$ ($c = 0$) $1.5 < t = 0.7 + 0.6 + 0.4 = 1.7$ ($c = 1$)

(c) Check termination

Figure 3: Intermediate variables for the execution of NTA for the example query. The values when $c=0$ are shown on the left, and the values when $c=1$ are shown on the right.

computed is $O(nNeurons \cdot nInputs \cdot \log_2(nInputs))$, but the main source of overhead is DNN inference, which is a cost proportional to the number of inputs. The method that DeepEverest adopts is more space-efficient than building an index over (neuronID, PID) pairs because it costs $nNeurons \cdot nInputs \cdot \log_2(nPartitions)$ bits rather than $nNeurons \cdot nInputs \cdot \log_2(nInputs)$ bits, where $nPartitions \ll nInputs$. NPI also has much smaller storage overhead compared to fully materializing all activation values. A PID takes less storage than an activation value because a PID only costs $\log_2(nPartitions)$ bits, while an activation value is usually a 32-bit floating point. For example, if DeepEverest has 8 partitions for each neuron, representing a PID costs 3 bits, which is less than 10% of the storage cost of full materialization. Storing the lower and upper bounds costs $nNeurons \cdot nPartitions \cdot 2 \cdot 32$ bits, which is normally negligible compared to the cost of storing the PIDs.

4.4 Neural Threshold Algorithm (NTA)

Notation. We denote with N the set of all neurons in the DNN and with D the input dataset. A neuron is denoted with $n \in N$ and an input with $x \in D$. x is the inputID. The user issues a query: $topk(s, G, k, DIST)$, where $s \in D$ is the sample input (also known as the target input) of interest to the user. $G \subseteq N$ is a set of neurons from

Table 2: Summary of frequently used notation.

Symbol	Meaning
s	Sample input (or target input) from D
G	Group of neurons from N
$topk(s, G, k, DIST)$	<i>Top-k most-similar</i> query
$DIST$	Function to compute distances between inputs
g_i (or i when clear)	The i -th neuron in G
$act(i, x)$	Activation value of g_i for input x
$dist(s, x, G)$	Distance between s and x based on G
top	Set of top- k inputs that are closest to s
P_n	Set of partitions for neuron n
$getInputIDs(n, p)$	inputIDs of a single partition $p \in P_n$
$sPID(n)$	PID to which s belongs for neuron n
$lBnd(n, p)$	Lower bound of a single partition $p \in P_n$
$uBnd(n, p)$	Upper bound of a single partition $p \in P_n$

a single layer in N . k is the desired number of query results, and $DIST$ is the distance function. This function computes the distance between the set of activations of s and x looking only at the neurons in G . Table 2 lists our frequently used notation.

NTA returns the set of top- k inputs that are closest to the target input when considering only the neurons in G . This set of top- k inputs can be defined as a set $top \subseteq D$ of k inputs. top is initially empty and is conceptually built incrementally by identifying and adding to top the next input that satisfies:

$$\arg \min_{x \in D \setminus top} \{dist(s, x, G)\} \quad (1)$$

We further denote with g_i (or i when clear) the i -th neuron in set G , and with $act(i, x)$ the activation of neuron g_i on input x . g_i is the neuronID. For each neuron $n \in N$, NPI includes the set of partitions, P_n . We denote a single partition for neuron n with $p \in P_n$, and p is the PID. We denote the lower and upper bounds of this partition $p \in P_n$ with $lBnd(n, p)$ and $uBnd(n, p)$.

NTA proceeds as follows,

Step 1: Load indexes. We assume that NPI is initially on disk. This step reads the NPI for the neurons $g_i \in G$ from disk. The index holds the set of partitions, their lower and upper bounds, and the PIDs of each input for each $g_i \in G$.

Step 2: Compute target activations. For each $g_i \in G$, compute $act(i, s)$, the activation value for input s and neuron g_i by running DNN inference on input s . A single inference pass is sufficient to compute the activations for all neurons in G .

Step 3: Order partitions. This step computes the order by which the partitions are accessed by NTA for each neuron. Let $sPID(i)$ denote the PID to which s belongs for neuron g_i . For each neuron $g_i \in G$ and partition $p \in P_i$, compute $dPar(i, p)$ as,

$$dPar(i, p) = \begin{cases} 0, & p = sPID(i) \\ lBnd(i, p) - act(i, s), & p < sPID(i) \\ act(i, s) - uBnd(i, p), & p > sPID(i) \end{cases} \quad (2)$$

which is the distance between the target input's activation value for neuron g_i and the closest activation value in partition p . For each neuron g_i , sort the partitions in P_i on their $dPar(i, p)$ values in ascending order and put them in a list, denoted with $ord(i)$. Later steps will process the partitions in the order specified by $ord(i)$.

Example: To illustrate the first three steps, consider a query $topk(x_5, \{R1, R2, R3\}, 2, l1\text{-distance})$ that finds the top-2 most

similar inputs to x_5 based on the activations of $\{R_1, R_2, R_3\}$, using the example dataset in Figure 1. In this example, $s=x_5$, $G=\{R_1, R_2, R_3\}$. Step 1 reads from disk PID , $lBnd$, and $uBnd$ shown in Figure 1. Step 2 runs DNN inference to compute the activations for x_5 , $(\text{act}(R_1, x_5), \text{act}(R_2, x_5), \text{act}(R_3, x_5))=(1.1, 1.1, 1.2)$. Step 3 computes $dPar$ and ord for $\{R_1, R_2, R_3\}$, as shown in Figure 2.

Step 4: Find top-k. This step runs the modified threshold algorithm. It starts with the partitions to which the target input belongs, and it expands its search from there. Unlike CTA, this step incrementally computes the activations for candidate inputs and does so in batches to get good GPU performance. This step proceeds as follows,

Starting with an index $c = 0$:

Step 4 (a): For each neuron g_i , maintain a set $toRun_i$ that contains the inputs whose activations should be computed. Access $ord(i, c)$ to get the partition that contains the next most similar inputs. Query NPI to get the inputIDs that belong to this partition $ord(i, c)$ and add them to $toRun_i$, i.e., $toRun_i \leftarrow \text{getInputIDs}(i, ord(i, c))$.

Example: as shown in Figure 3a, when $c = 0$, $toRun_{R_1, 0} = \{x_4, x_5\}$ because $ord(R_1, 0) = 2$.

Step 4 (b): For each neuron g_i , compute the activations for the inputs in $toRun_i$ (excluding those that have already been computed) by running DNN inference in batches. $toRun_i$ is cleared after DNN inference. Note that this inference step computes the activations for *all* neurons in the neuron group being queried. Compute the distance between each newly computed input x and the target input s as $dist(s, x, G) = \text{DIST}(|\text{act}(0, x) - \text{act}(0, s)|, \dots, |\text{act}(|G|-1, x) - \text{act}(|G|-1, s)|)$. Update top if $dist(s, x, G)$ is one of the k -smallest NTA has seen so far, i.e., input x is one of the k -most similar inputs to the target input s seen so far. Ties are broken arbitrarily.

Example: as shown in Figure 3b, when $c=0$, the activations of inputs x_2, x_4 are computed (x_5 was computed in Step 1). The distances from x_2 and x_4 to x_5 are 1.5 and 0.3, respectively.

Step 4 (c): Maintain a range of seen activations for inputs from $toRun_i$ for each neuron g_i , which is the range of activations such that NTA has seen every input with an activation in the open interval of this range. It is possible that NTA has seen one or more inputs from other neurons' $toRun$ sets with activations outside of this range. However, the open interval of this range denoted by $(minBoundary_i, maxBoundary_i)$ only contains the activations for the inputs that NTA is guaranteed to have seen.

Let $minDist_i$ be the shorter distance from the boundaries of this range to the target input for each neuron g_i : $minDist_i = \min\{F_i \cdot |minBoundary_i - \text{act}(i, s)|, V_i \cdot |maxBoundary_i - \text{act}(i, s)|\}$, where F_i is an indicator function that indicates whether NTA has seen the last partition (inputs with lowest activations) of neuron g_i , and V_i is another indicator function that indicates whether the 0-th partition (inputs with highest activations) of neuron g_i has been seen. Specifically, $F_i = \infty$ when the last partition of neuron g_i has been seen; $F_i = 1$ otherwise. $V_i = \infty$ when the first partition of neuron g_i has been seen; $V_i = 1$ otherwise. Define the threshold to be,

$$t = \text{DIST}(minDist_0, minDist_1, \dots, minDist_{|G|-1}) \quad (3)$$

The threshold, t , represents the smallest possible distance to s from any unseen input. Hence, the termination condition is,

$$\max_{x \in top} \{dist(s, x, G)\} \leq t \quad (4)$$

where $\max_{x \in top} \{dist(s, x, G)\}$ represents the maximum distance to the target input s in the current top- k result set. As soon as this inequality holds, halt and return top as the query results.

Example: as shown in Figure 3c, $minBoundary_i$, $maxBoundary_i$ and $minDist_i$ are maintained and calculated for $\{R_1, R_2, R_3\}$. For example, when $c = 0$, $minBoundary_{R_1} = 1.1$, $maxBoundary_{R_1} = 1.2$. Since NTA has seen the last partition (2) and has not seen the first partition (0), $F_{R_1} = \infty$, $V_{R_1} = 1$. Therefore, $minDist_{R_1} = |maxBoundary_{R_1} - \text{act}_{R_1, x_5}| = |1.2 - 1.1| = 0.1$.

When $c = 0$, $t = 0.2 < 1.5 = \max_{x \in top} \{dist(s, x, G)\}$, so NTA does not halt. When $c = 1$, $t = 1.7 \geq 1.5 = \max_{x \in top} \{dist(s, x, G)\}$, so NTA halts and returns top as the query result. It is worth noting that the cost of DNN inference on x_0 is not incurred because it is impossible for x_0 to be one of the top-2 results.

Step 4 (d): Increment c by 1. Repeat Step 4 (a) - (d) until all partitions have been seen or the halting condition in Step 4 (c) is satisfied.

The key innovation of NTA compared to CTA is in its processing of the inputs partition-by-partition using NPI until the termination condition is met. NTA runs DNN inference on only the necessary partitions of inputs for it to be certain that it has the precise top- k results when it terminates. This approach significantly reduces the number of inputs on which DNN inference is performed at query time compared to computing the activation values for all inputs at query time. It further improves query performance by utilizing batch processing on GPUs [40]. Inputs that share a partition are sent to the DNN for inference all at once. NTA along with NPI also has much smaller storage overhead compared to fully materializing the activations for all inputs. We include NTA's pseudocode and discussions of approximation, early stopping, and incrementally returning query results in [17] due to space constraints.

4.5 Instance Optimality of NTA

In this section, we investigate the instance optimality of NTA, which corresponds to the optimality in every instance, rather than just the worst or average case. Fagin's paper [11] shows that CTA is instance optimal for finding the top- k items over all algorithms (excluding those that make very lucky guesses) and over all legal databases. We build on that proof to show the same for NTA.

Following the definitions in [11], let \mathbb{A} be a set of algorithms that answer our target queries, and \mathbb{D} be a set of combinations of datasets and DNN models that are legal inputs. Let $cost(\mathcal{A}, \mathcal{D})$ be the number of inputs in the dataset of \mathcal{D} accessed by $\mathcal{A} \in \mathbb{A}$ when running \mathcal{A} on $\mathcal{D} \in \mathbb{D}$. An algorithm \mathcal{B} is instance optimal over \mathbb{A} and \mathbb{D} if $\mathcal{B} \in \mathbb{A}$ and if for every $\mathcal{A} \in \mathbb{A}$ and every $\mathcal{D} \in \mathbb{D}$,

$$cost(\mathcal{B}, \mathcal{D}) = O(cost(\mathcal{A}, \mathcal{D})) \quad (5)$$

We show the following theorem.

THEOREM 4.1. *NTA is instance optimal for finding the top- k inputs over all algorithms (excluding those that make very lucky guesses) and over all legal combinations of datasets and DNN models.*

The proof follows similarly to that of Theorem 6.1 in [11]. We bound the maximal number of inputs accessed by NTA with an additive constant over CTA's maximal sequential access depth.

PROOF. For any $\mathcal{D} \in \mathbb{D}$, assume that we already have the relation of sorted absolute differences (denoted with $AbsDiff$) between the activations of all inputs and the activation of the target input. Assume that CTA halts at depth d_i for each neuron g_i when running

on AbsDiff, i.e., d_i is the number of inputs seen via sequential accesses for neuron g_i . As in [11], it suffices to bound the maximal number of inputs accessed by NTA for any $g_i \in G$. Let $d = \max_i d_i$. For the duration of this proof, let x_j denote the input at depth j in AbsDiff for a neuron g_i that reaches depth d when running CTA. Let R be the partition size in NPI. We will show that NTA will have accessed $x_j (\forall 1 \leq j \leq d)$ and terminate in at most $d + 2R$ accesses.

Recall that we denote a single partition for neuron g_i with $p \in P_i$. p is the partition identifier, PID, and $\text{sPID}(i)$ is the PID of the target input s for neuron g_i . In NPI, for $g_i \in G$, we say that a partition $p \in P_i$ is an *above* partition if $p < \text{sPID}(i)$; a partition $p \in P_i$ is an *under* partition if $p \geq \text{sPID}(i)$. We say that an input is an *above* input if it belongs to an *above* partition; an input is an *under* input if it belongs to an *under* partition. As in eq. (2), an *above* partition uses the input whose activation is the lower bound of the partition as its representative; an *under* partition uses the input whose activation is the upper bound as its representative (the only exception is that the partition $p = \text{sPID}(i)$ uses the target input, s , as its representative).

Without loss of generality, assume that x_d is an *above* input. Let a be the greatest $j < d$ where x_j is an *under* input. Let b be the least $j > d$ where x_j is an *under* input. When NTA has accessed $x_j (\forall 1 \leq j \leq d)$ and confirmed that x_b is more distant from s than x_d , it knows that it has the correct top- k results and can then terminate. This is equivalent to the termination condition in Section 4.4.

Case 1: When x_a is accessed by NTA after x_d , we will show at most R *above* inputs, $x_j (j > d)$, are accessed before x_a . Assume towards contradiction that more than R *above* inputs $x_j (j > d)$ are accessed before x_a . There would be a representative of an *above* partition, $x_{d'}$ where $d' \geq d$. Let $x_{a'}$ be the representative of the partition containing x_a . Thus, $a' \leq a < d \leq d' \implies a' < d'$. This is a contradiction because the *above* partition with $x_{d'}$ as its representative was chosen to be accessed earlier than the *under* partition with $x_{a'}$ as its representative, which implies $d' \leq a'$.

Since *under* partitions are accessed in order w.r.t. x_a , the number of *under* inputs, $x_j (j > d)$, accessed by NTA is also at most R , i.e., the *under* inputs co-located on the partition containing x_a .

Note that if x_a and x_b belong to the same partition, NTA will terminate after this partition. If x_a and x_b belong to different partitions, then x_b is the representative of its partition. After confirming that x_b is more distant from the target input s than x_d , NTA knows that it does not need to access any further partitions (including the partition containing x_b) and terminates. In either scenario, the number of *above* $x_j (j > d)$ accessed by NTA is at most R , and the number of *under* $x_j (j > d)$ accessed is also at most R . Hence, the total number of accesses made by NTA is at most $d + 2R$.

Case 2: When x_b is accessed by NTA before x_d , there can be at most R *under* inputs, $x_j (j > d)$, that are accessed before x_d . Furthermore, the number of *above* inputs, $x_j (j > d)$, accessed is at most R , i.e., the *above* inputs co-located on the partition containing x_d . Hence, the total number of accesses made by NTA is at most $d + 2R$. The proof follows similarly to Case 1.

Case 3: When x_a, x_b are accessed by NTA in order w.r.t. x_d, x_b must be the representative for its partition. After NTA processes the partition containing x_d , it will recognize that x_b is more distant than x_d and terminate. Thus no *under* $x_j (j > d)$ will be explicitly accessed. The *above* partitions are accessed in order w.r.t. x_d , so the number of *above* inputs, $x_j (j > d)$, accessed is at most R , i.e., the

inputs co-located on the partition containing x_d . Hence, the total number of accesses made by NTA is at most $d + R$.

Since the three cases above exhaust all possibilities, this proves that for each neuron in G , NTA will have accessed $x_j (\forall 1 \leq j \leq d)$ and terminate in at most $d + 2R$ accesses. \square

4.6 Incremental Indexing

As shown in Section 5, the DeepEverest approach described so far achieves excellent query execution times with low storage overhead. This approach, however, incurs a potentially high preprocessing cost, especially for large datasets and models. Before executing any query, DeepEverest needs to compute the activations for all neurons and all inputs by running DNN inference. It then needs to construct the indexes for all layers and persist them to disk.

To address this challenge, we propose building the indexes incrementally as queries execute so that preprocessing is performed *only for the layers users query*. With this approach, DeepEverest performs no preprocessing ahead of time. When the user submits a query, if the indexes of the queried layer are available on disk, DeepEverest proceeds as described in Sections 4.4 and 4.7.1; otherwise, DeepEverest computes the activations of the queried layer by running DNN inference on all inputs. While doing so, it computes the query answer and returns it to the user. DeepEverest then constructs the indexes for the layer and persists them to disk. Note that DNN inference is performed starting from the first layer instead of a previously queried layer each time DeepEverest constructs the indexes for a layer because only the indexes of the queried layers are stored on disk. With this approach, the costs of index computation and persistence for each layer are incurred once the first time that layer is queried, and only if that layer is queried.

In Section 5.3, we show that DeepEverest with this incremental approach significantly outperforms other methods on multi-query workloads. While DeepEverest must do extra preprocessing to build and store its indexes compared with caching the activations directly, it accelerates significantly more queries because it is able to store the indexes for significantly more layers given a storage budget.

4.7 Optimizations

In this section, we present several important optimizations that further improve the performance of DeepEverest. The first optimization, described in Section 4.7.1, accelerates two common types of top- k queries. The second optimization, described in Section 4.7.2, automatically selects DeepEverest’s configuration. The third optimization, described in Section 4.7.3, accelerates sequences of related queries, as may occur during data exploration.

4.7.1 Maximum Activation Index (MAI). For a given target input and a layer in a DNN, the *maximally activated neurons* are those neurons in the layer for which the activation values for the target input are the highest. DNN interpretation often involves examining such maximally activated neurons [5, 50, 59, 62] because they respond to the input the most and have the greatest impact on the DNN output. A common set of *top- k most-similar* queries ask to find the top- k similar inputs to a target input based on a neuron group consisting of these maximally activated neurons.

To accelerate these *top- k most-similar* queries that target maximally activated neurons as well as *top- k highest* queries, we introduce a straightforward yet effective optimization. The idea is

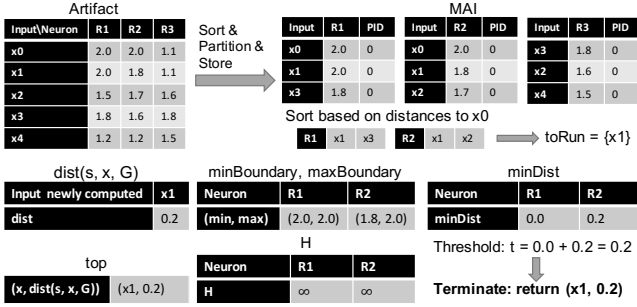


Figure 4: An example of constructing MAI ($ratio=0.6$) and query execution for $topk(x0, \{R1, R2, R3\}, 1, l1\text{-distance})$ ($batchSize=1$). Despite $x0$ only being in MAI for R1 and R2, DeepEverest leverages MAI to answer the query after only running DNN inference on $x0$ and $x1$.

for DeepEverest to store, for each neuron, a fraction of the highest activation values together with the corresponding inputIDs. We call this data structure the Maximum Activation Index (MAI), and denote this fraction of (activation, inputID) pairs for a given neuronID with $MAI(neuronID)$. This fraction automatically becomes each neuron’s 0-th partition. We denote the fraction of the inputs stored in MAI with $ratio$, which is a configurable parameter and discussed further in Section 4.7.2.

DeepEverest utilizes MAI during query execution to further reduce the number of inputs on which DNN inference is performed, if possible. DeepEverest now has more detailed knowledge of which inputs are most similar to the target input, rather than just the high-level knowledge that the inputs are in the same partition. We observe empirically that the activation values of the *maximally activated neurons* for an input are often likely to be in the top activations stored in MAI, and thus MAI is effective in improving the query time. DeepEverest modifies query execution described in Section 4.4 of partition 0 to incorporate this information as follows: it first finds the neurons for which the target input is in MAI. For these neurons, DeepEverest sorts the other inputs in the 0-th partition by their distances to the target input s . Rather than performing DNN inference on all inputs in MAI for each neuron, DeepEverest builds a global *toRun* set by adding the most similar inputs from all of these neurons until the batch size is reached. Step 4(c) in Section 4.4 is modified to compute $minDist_i$ as $\min\{|minBoundary_i - act_{i,s}|, H_i \cdot |maxBoundary_i - act_{i,s}|\}$, where H_i is an indicator function that indicates whether NTA has seen the input with the highest activation in MAI (i). $H_i = \infty$ when highest activation of neuron g_i has been seen; $H_i = 1$ otherwise. The neurons g_i for which s is not in MAI(i) contribute 0 to the threshold calculation. Figure 4 illustrates an example.

4.7.2 Automatic Configuration Selection. Given a storage budget, DeepEverest must allocate it between NPI and MAI. It uses a heuristic algorithm to achieve this. A greater $nPartitions$ leads to smaller partitions, which is key to generally better performance (see Section 5.4) on queries that target any kind of a neuron group (see Section 5.1), while a larger $ratio$ accelerates only the two types of queries mentioned in Section 4.7.1. Hence, DeepEverest first picks a value for $nPartitions$ and then sets the value of $ratio$.

Intuitively when partitions are smaller, DNN inference is performed on fewer inputs not in the top- k because DeepEverest processes inputs partition-by-partition. However, if the partitions are smaller than the optimal batch size, DeepEverest will not leverage the full GPU parallelism.

Given a storage budget, $budget$ (in bytes), and a batch size, $batchSize$, DeepEverest sets $nPartitions$ to be the maximum power of two (to utilize all bits) that satisfies $nPartitions \leq nInputs/batchSize$ and $cost(nPartitions) < budget$. $cost(nPartitions)$ is the bytes consumed by storing NPI and is calculated as $nNeurons \cdot nInputs \cdot \log_2(nPartitions)/8$. $batchSize$ is set to the value that achieves the highest throughput for the DNN. Given the remaining storage budget, DeepEverest sets $ratio$ to be the maximum value that satisfies $cost(ratio) \leq budget - cost(nPartitions)$, where $cost(ratio)$ is the bytes consumed by storing MAI and calculated as $ratio \cdot nInputs \cdot nNeurons \cdot 4 \cdot 2$, since activation and inputID are 4 bytes each. When there is no remaining storage budget after selecting $nPartitions$, DeepEverest sets $ratio$ to 0.

4.7.3 Inter-Query Acceleration (IQA). Inter-Query Acceleration (IQA) is an optimization technique to accelerate sequences of related queries, as may occur during DNN interpretation. As an example, imagine that a user finds a misclassified image. The user may want to first see the maximally activated neurons in a layer for the image and then find images with similar maximally activated neurons. The user may then decide to change how many neurons they are looking at, e.g., go from the top-3 neurons to the top-4 neurons. Queries that overlap in neurons present an opportunity for further optimization as activation values can be reused for related queries.

With IQA, DeepEverest leverages an in-memory cache that contains recently used activation values to reduce the number of inputs that it must run DNN inference on at query time. Note that this in-memory cache is different from the disk caches described in Section 4.1. During query execution, DeepEverest inserts the activation values of each input processed by NTA into the cache. Instead of caching only the activation values for the neuron group being queried, it caches the activations for *all* neurons in the queried layer. This enables DeepEverest to utilize the cache for future related queries that target a different group of neurons in the same layer. DeepEverest adopts a most recently used (MRU) replacement policy for the in-memory cache. This is because DeepEverest processes partitions in order from most similar to the target input to least similar, and it seeks to prioritize keeping the activations from the most similar partitions in the cache. We show in Section 5.6 that given a small in-memory cache budget, DeepEverest with IQA achieves up to 8 \times faster query times than DeepEverest without it.

5 EVALUATION

DeepEverest is implemented in Python, using C++ to build the indexes. We evaluate it against the baselines described in Section 4.1.

5.1 Evaluation Setup

Datasets and models. We evaluate DeepEverest on two sets of well-known datasets and models. The first, called *CIFAR10-VGG16*, uses as inputs 10,000 images from the test set of CIFAR10 [27], and uses a VGG16 network [31, 48]. The second, called *ImageNet-ResNet50*, uses as inputs 10,000 images from the validation set of

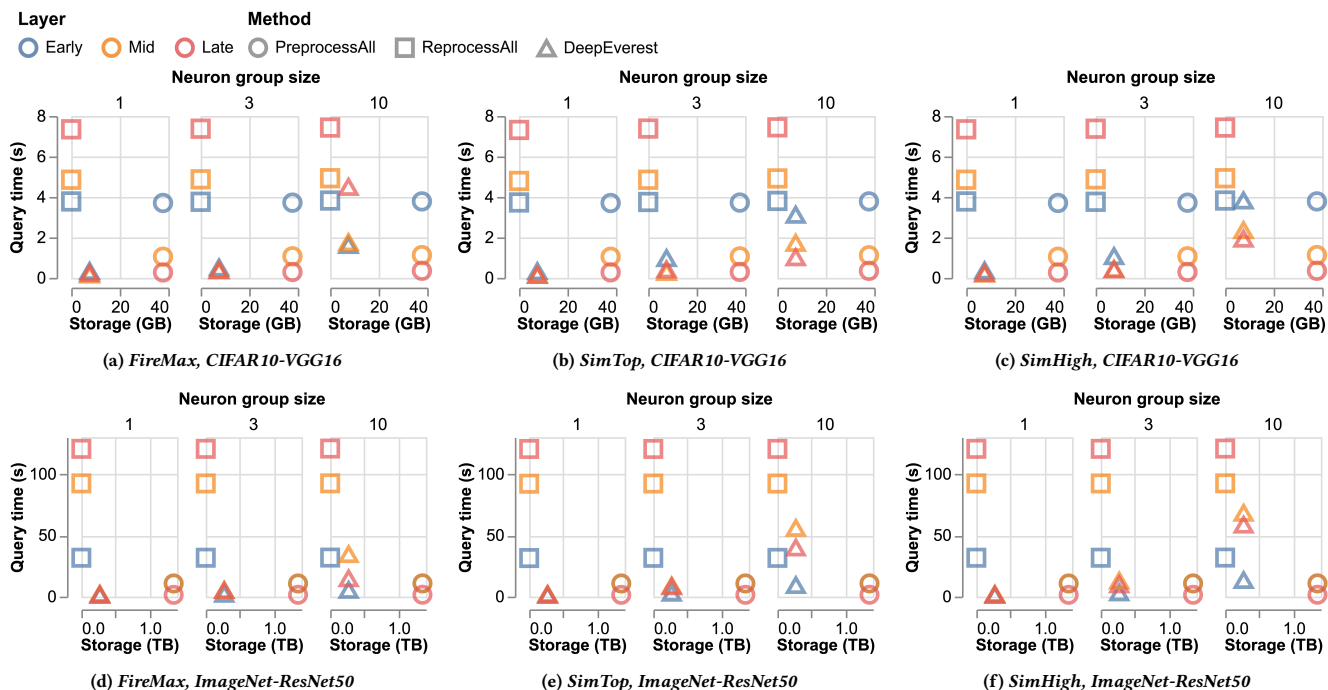


Figure 5: End-to-end individual query times and storage sizes on *CIFAR10-VGG16* and *ImageNet-ResNet50*. $nPartitions$ and ratio of DeepEverest are selected by our heuristic algorithm given a storage budget of 20% of full materialization.

ImageNet [46], and uses a ResNet50 model [18]. These two sets of models and datasets complement each other in terms of model and input size and DNN inference cost. In all experiments, we pre-load the entire input dataset into memory. We set $batchSize$ for each model as the value that achieves the highest inference throughput (128 for *CIFAR10-VGG16*; 64 for *ImageNet-ResNet50*).

Query generation. To generate queries, we consider 3 types of layers: *early*, *mid*, and *late*. For *CIFAR10-VGG16*, these correspond to activation layers 2, 7, and 13. For *ImageNet-ResNet50*, we use activation layers 2, 25, and 48. Given an input and a layer, we consider the following types of neuron groups: (a) *Top*: the maximally activated neurons for the given input in the layer; and (b) *RandHigh*: neurons randomly picked from the top half of non-zero neurons for the given input. We further consider *small*, *medium*, and *large* neuron groups consisting of 1, 3, and 10 neurons. Finally, based on the neuron groups, we use the following query types: (a) *FireMax*: *top-k* highest query; (b) *SimTop*: *top-k* most-similar query based on a *Top* neuron group; and (c) *SimHigh*: *top-k* most-similar query based on a *RandHigh* neuron group. We randomly select inputs from each dataset to generate *SimTop* and *SimHigh* queries.

In all experiments, we set $k=20$, which is a reasonable number of results for a user to inspect for a query. We use l_2 -distance as the distance function. All numbers reported are median values of five queries on random inputs for each query configuration (e.g., query type: *FireMax*, neuron group size: 3, layer: *late*).

Machine configuration. All experiments are run on an AWS EC2 p2.xlarge instance, which has an Intel Xeon E5-2686 v4 CPU running at 2.3 GHz, with 61 GB of RAM, an NVIDIA K80 GPU with 12 GB of GPU memory, and EBS gp3 volumes for disk storage.

5.2 Fundamental Space-Time Tradeoff

We first evaluate the fundamental trade-off that DeepEverest achieves in terms of storage space and query execution time for individual queries. In this experiment, the only optimization DeepEverest uses is MAI described in Section 4.7.1. We first precompute and store the indexes for all layers before executing the benchmark queries. DeepEverest has a storage budget of 20% of *PreprocessAll*, and selects $nPartitions$ and $ratio$ using the algorithm described in Section 4.7.2. For *CIFAR10-VGG16*, $nPartitions = 64$, $ratio = 0.0046$; for *ImageNet-ResNet50*, $nPartitions = 64$, $ratio = 0.0074$. We compare DeepEverest against *PreprocessAll* and *ReprocessAll*. Figure 5 shows the results.

As the figures show, *PreprocessAll* has the highest storage cost (37.8 GB for *CIFAR10-VGG16*, and 1.35 TB for *ImageNet-ResNet50*) since it stores all activations for every input. However, scanning the precomputed activation values generally leads to the fastest query times. The query times of *PreprocessAll* are slower for the early layer of *CIFAR10-VGG16* because it has a large number of neurons, and thus it takes longer to load all the activations. *ReprocessAll* has the lowest storage cost since it does not precompute or store anything ahead of time. Its query times are slow because of the DNN inference on the entire dataset at query time.

DeepEverest achieves the best of both worlds: low storage overhead and fast query times. For *CIFAR10-VGG16*, compared with *ReprocessAll* DeepEverest is $1.65\times$ to $31.1\times$ faster for *FireMax*, $1.22\times$ to $50.8\times$ faster for *SimTop*, and up to $31.5\times$ faster for *SimHigh*. For *ImageNet-ResNet50*, compared with *ReprocessAll*, DeepEverest is $2.67\times$ to $62.8\times$ faster for *FireMax*, $1.65\times$ to $63.5\times$ faster for *SimTop*, and $1.35\times$ to $63.1\times$ faster for *SimHigh*.

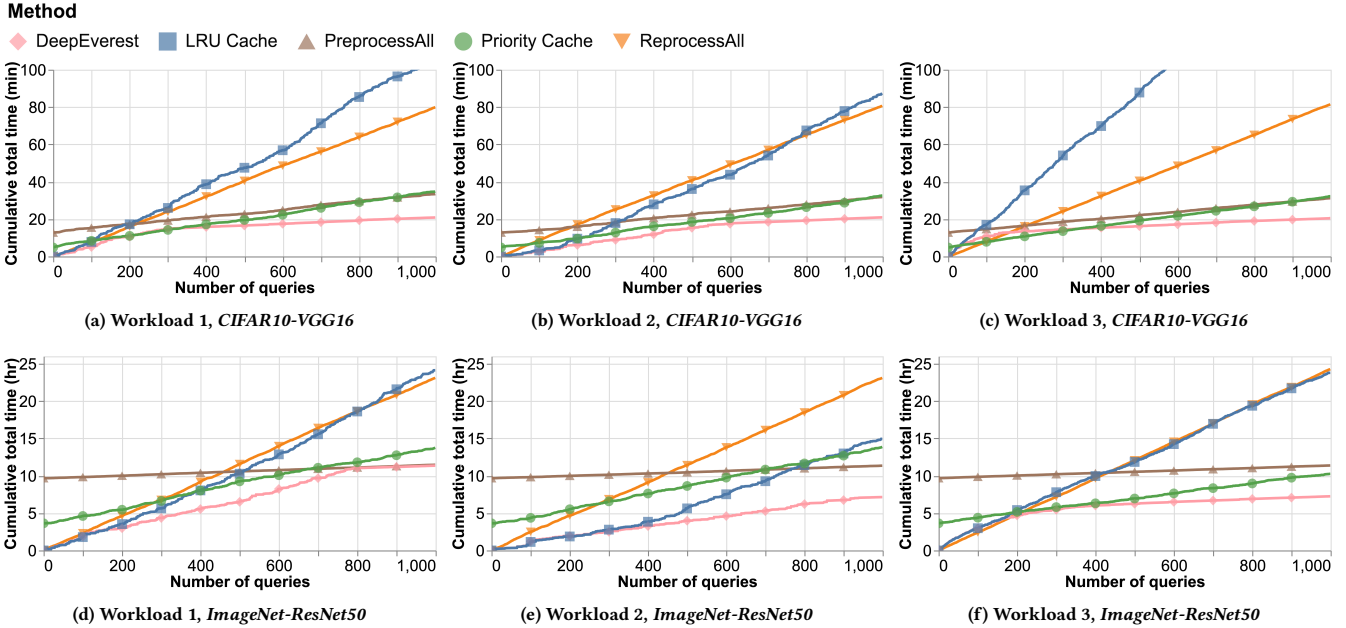


Figure 6: Cumulative total time (preprocessing time plus query execution time) for various multi-query workloads.

Compared to *PreprocessAll* for both *CIFAR10-VGG16* and *ImageNet-ResNet50*, *DeepEverest* achieves comparable and sometimes even faster query times for queries that target small and medium-size neuron groups despite using only 20% of *PreprocessAll*'s storage overhead. For queries that target large neuron groups, *DeepEverest*'s query times are slower. Due to the curse of dimensionality, there is little difference in the distances between different pairs of inputs. As a result, *DeepEverest* is not able to reduce the number of inputs run by the DNN at query time as it does for small and medium neuron groups. Table 3 shows the number of inputs run by the DNN at query time to compute the activation values for *SimHigh* queries. We find that the number of inputs run by the DNN at query time for queries on larger neuron groups is higher than that of queries on smaller neuron groups.

5.3 Multi-Query Workloads

In this section, we evaluate *DeepEverest* on multi-query workloads using incremental indexing described in Section 4.6 that avoids start-up overhead. We construct various query workloads to represent possible DNN interpretation patterns and compare *DeepEverest* against other on-disk caching techniques. All workloads consist of 1,000 the most general *SimHigh* queries that target neuron groups of medium size. The first query of each workload targets a *RandHigh* neuron group from a randomly selected layer. Each later query has probabilities p_{same} of querying the same layer as the previous query, p_{prev} to query one of the previously queried layers (excluding the layer queried by the previous query), and p_{new} to query a layer that has not been queried yet. Workload 1 sets these to $p_{same}=0.5, p_{prev}=0.3, p_{new}=0.2$. Workload 2 sets these to $p_{same}=0.5, p_{prev}=0.4, p_{new}=0.1$. Workloads 1 and 2 are intended to simulate the exploration process of users that are likely to initially target layers they are interested in, and gradually explore more

layers. Additionally, we construct Workload 3 in which queries are independent of each other; each layer is targeted uniformly at random by each query. This is not a realistic interpretation pattern but is meant to show the worst-case workload for *DeepEverest*.

We measure the cumulative total time, which includes the time for both preprocessing and query execution, and cumulative storage for each method. *DeepEverest* is given a storage budget of 20% of full materialization, and $nPartitions$ and $ratio$ are selected by our heuristic algorithm. *LRU Cache* and *Priority Cache* have the same 20% storage budget. The time to initially compute and store the data on disk is included with the 0-th query for *PreprocessAll* and *Priority Cache*. The results for cumulative total time are shown in Figure 6. We report the storage results in the text.

DeepEverest consistently performs the best for Workloads 1 and 2 using less than 20% of the storage of full materialization. We observe that after some number of queries, the cumulative total time of *DeepEverest* grows more slowly. For *CIFAR10-VGG16*, it plateaus after around 300 queries for Workload 1 and around 550 queries for Workload 2. This indicates that *DeepEverest* has built and stored NPI and MAI for all layers in the DNN. All later queries are much faster because they benefit from these indexes and NTA. For *ImageNet-ResNet50*, *DeepEverest* completes building and storing the indexes for all layers after around 780 queries for Workload 1 and never completes for Workload 2, as we observe that *DeepEverest*'s storage consumption is only 11.3% of full materialization after 1,000 queries. *DeepEverest* finishes building its indexes after fewer queries in Workload 1 than in Workload 2 since Workload 1 has a higher probability of querying new layers.

While *DeepEverest* has the fastest query times, its storage also grows more slowly than the baseline approaches (except for *ReprocessAll* which does not have any storage overhead). For both datasets and models, *PreprocessAll* uses full storage after

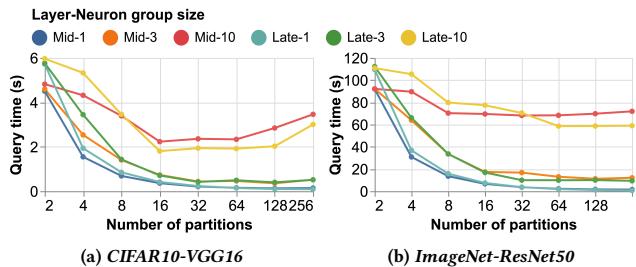


Figure 7: Query times of *SimHigh* queries when varying $nPartitions$. Note the log scale on the x -axis.

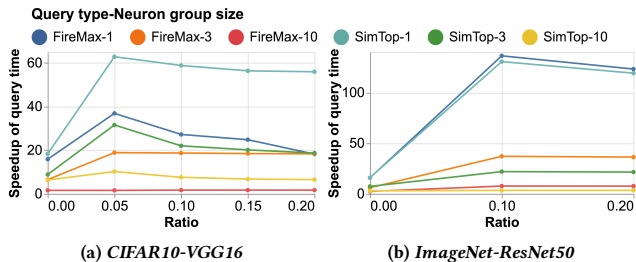


Figure 8: Speedups of query times (layer: *late*) against *ReprocessAll* when varying $ratio$ with $nPartitions$ set to 16.

its preprocessing step. Similarly, *Priority Cache* consumes its 20% storage budget after preprocessing. *LRU Cache* consumes its storage budget after around 50 to 200 queries. As discussed above, *DeepEverest* finishes building the indexes for all layers and consumes its storage budget after around 300 to 500 queries on *CIFAR10-VGG16*, and for *ImageNet-ResNet50* it fills its storage after 780 queries for Workload 1 and never does so for Workload 2.

For Workload 3, which is unlikely an interpretation pattern, *DeepEverest* is slightly worse than the best performing method for the first 200 to 300 queries on both datasets and models because during that time *DeepEverest* builds indexes for many new layers that have not been queried before. However, *DeepEverest* performs the best after 400 queries because more queries target previously seen layers and benefit from its indexes and NTA.

We further observe that users typically pause between queries. *DeepEverest* can use that time to compute and persist its indexes to disk, which would yield even better user-perceived query times.

5.4 *DeepEverest*'s Configuration Selection

We now study the effectiveness of NPI and MAI, and the impact of *DeepEverest*'s configurable parameters, as well as how *DeepEverest* performs using the configuration selection heuristic described in Section 4.7.2 with different storage budgets.

Impact of Number of Partitions. We first examine how the number of partitions, $nPartitions$, affects the query times. We measure the query times of *DeepEverest* on *SimHigh* queries after building NPI with varying $nPartitions$. MAI is disabled for this experiment. The results are shown in Figure 7. We also measure the number of inputs on which *DeepEverest* performs DNN inference during query processing. Table 3 shows the results for *CIFAR10-VGG16*. Similar trends are observed for *ImageNet-ResNet50*.

The query time initially decreases as $nPartitions$ increases. This is because when partitions are larger, inputs that do not contribute

Table 3: Number of inputs run by the DNN at query time for *SimHigh* queries on *CIFAR10-VGG16*.

Layer-Neuron group size	Number of partitions						
	4	8	16	32	64	128	256
mid-1	3334	1429	667	323	159	79	40
mid-3	5462	2902	1441	736	727	390	390
mid-10	8941	6869	4339	4215	3515	3492	3316
late-1	3334	1429	667	323	159	79	40
late-3	5968	2372	1106	618	618	388	391
late-10	9008	5565	2870	2745	2227	1956	1919

to the result end up being processed by *DeepEverest*. Hence, as partitions get smaller, the number of inputs run by the DNN at query time decreases. Then, after $nPartitions$ increases past a certain value (64 for *CIFAR10-VGG16*; 128 for *ImageNet-ResNet50*), the query time no longer decreases despite the number of inputs run by the DNN at query time continuing to decrease. Recall that NTA runs inference on all inputs in a partition as it processes that partition. When $nPartitions$ is so large that the partition size is below the optimal *batchSize* for DNN inference, the parallelism of the GPU is not fully utilized, which causes some queries to slow down. Therefore, a good value of $nPartitions$ creates partitions whose sizes are similar to the optimal *batchSize*.

Effectiveness of MAI. This experiment evaluates the effectiveness of MAI. We measure the speedups of query times compared with *ReprocessAll* when varying $ratio$, which determines the fraction of inputs with activation values materialized in MAI. Recall that when MAI is non-empty, it becomes the 0-th partition. For this experiment, we set $nPartitions = 16$, which is a setting that performs well (see Figure 7). As discussed in Section 4.7.1, MAI is designed to accelerate *FireMax* and *SimTop* queries. We measure the speedups of such queries on neuron groups of different sizes.

Figure 8 shows the results. Note that when $ratio=0$, *DeepEverest* runs without MAI. The speedups of query times are generally much higher when $ratio$ is any non-zero value. This is because MAI enables *DeepEverest* to return the query results after processing a subset of the inputs from MAI (partition 0), rather than processing the entire partition. We also observe that the speedups of query times plateau or drop as $ratio$ further increases. This is because loading MAI from disk takes longer as $ratio$ increases. When a small index provides enough information for *DeepEverest* to find the top- k results after processing only some inputs from MAI, increasing $ratio$ degrades the speedups; the additional inputs in MAI do not improve the query times, and loading a larger index takes longer. The best value of $ratio$ in practice depends on the queries and the distributions of the activations of the queried neuron group. Empirically, we observe that a small value of $ratio$ (e.g., 0.05) is good for *FireMax* and *SimTop* queries on the two datasets and models.

Impact of Storage Budget. In previous sections, we examined the performance of *DeepEverest* with a storage budget of 20% of full materialization. Here we examine how well *DeepEverest* performs when the configuration selection algorithm has different storage budgets. We measure the speedups of query times compared with *ReprocessAll* for *SimTop* and *SimHigh* queries that target medium and large neuron groups, as shown in Figure 9. We

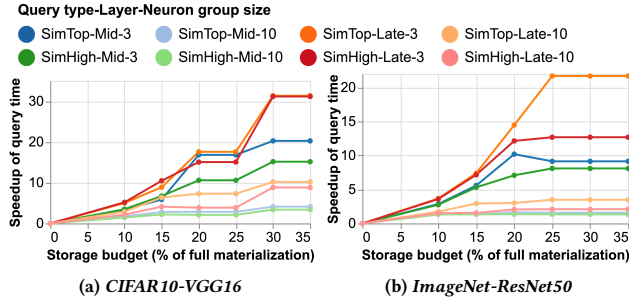


Figure 9: Speedups against *ReprocessAll* by DeepEverest when given different storage budgets.

observe that empirically DeepEverest delivers high speedups across different storage budgets, which also suggests that our configuration selection algorithm is robust. With a larger storage budget, DeepEverest performs better. We also observe that the speedups of queries on medium neuron groups are generally greater than the speedups for queries on large neuron groups.

5.5 Preprocessing Costs

This experiment evaluates the costs of preprocessing in an extreme case where the user queries all layers in the DNN. Note that with incremental indexing, the costs of index computation and persistence for each layer are incurred once *only* if that layer is queried. In this experiment, DeepEverest is given a storage budget of 20% of full materialization and preprocesses all layers for each dataset and model from the first layer to the last layer. Convolutional layers, activation layers, and batch normalization layers are considered separate layers. We measure the cumulative times for each component in preprocessing: DNN inference, data persistence, and index computation. We force-write the data to disk when measuring the time for data persistence. Figure 10 shows the results. DeepEverest has similar preprocessing times compared with *PreprocessAll*. The time for building NPI and MAI and persisting them to disk is similar to the time for *PreprocessAll* to persist the activations to disk. Considering these results along with the results shown in Section 5.2, DeepEverest achieves comparable and sometimes better query times than *PreprocessAll*, with only 20% of its storage overhead and similar preprocessing times.

5.6 Effectiveness of IQA

Finally, we evaluate the effectiveness of Inter-Query Acceleration (IQA) for sequences of related queries. We randomly select five inputs and construct two sequences of related queries for each input on various layers. Both sequences consist of 1,000 *SimHigh* queries. The first query of each sequence targets a *RandHigh* neuron group containing n_{size} neurons. Each later query randomly replaces $n_{replace}$ neurons in the neuron group of the previous query by $n_{replace}$ randomly selected *RandHigh* neurons. Sequence 1 sets $n_{size}=5$ and $n_{replace}=1$. Sequence 2 sets $n_{size}=10$ and $n_{replace}=2$.

We measure the speedups for query times of DeepEverest with IQA against DeepEverest without IQA for each query. Figure 11 shows the median of the speedups for each query on *CIFAR10-VGG16* given an in-memory cache budget of 1 GB for IQA, with $nPartitions=16$ and $ratio=0$ set for DeepEverest. We observe

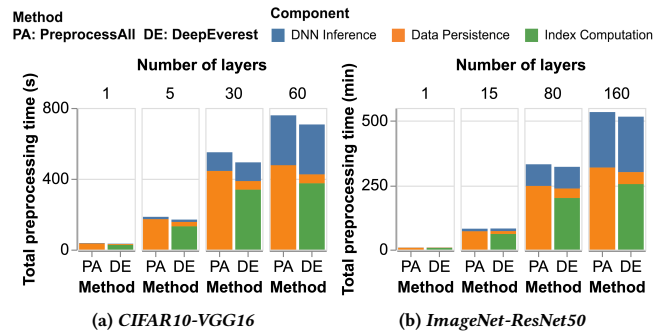


Figure 10: Cumulative preprocessing times from the first layer to the last layer for *PreprocessAll* and DeepEverest.

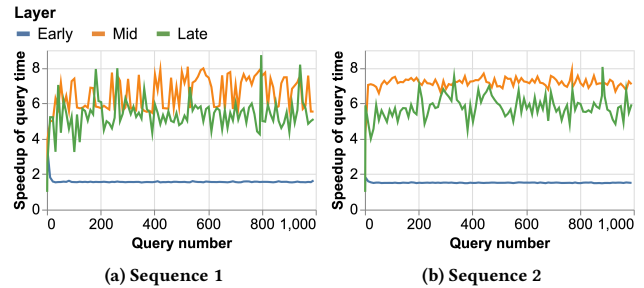


Figure 11: Speedups of query times by DeepEverest with IQA (1 GB cache budget) against DeepEverest without IQA.

that even with this small budget, IQA consistently improves DeepEverest’s query times across different layers. Not shown in the paper, we also experiment with different $nPartitions$ and $ratio$ and different cache budgets. We find that IQA always consistently speeds up related queries and larger budgets generally lead to larger speedups. In Figure 11, the speedups for the first query are around $1\times$ since the in-memory cache is initially empty. For later queries when the cache is populated, for Sequence 1, DeepEverest with IQA achieves speedups of 2.65 \times to 8.73 \times for the late layer, 3.97 \times to 8.08 \times for the mid layer, and 1.53 \times to 3.38 \times for the early layer. For Sequence 2, DeepEverest with IQA achieves speedups of 4.00 \times to 8.06 \times for the late layer, 4.29 \times to 7.86 \times for the mid layer, and 1.48 \times to 1.82 \times for the early layer. The speedups for the early layer are smaller because this layer is larger, and hence the in-memory cache can hold fewer inputs’ activations of the full layer. We observe larger speedups for the early layer with larger cache budgets.

6 CONCLUSION

We presented DeepEverest, a system that accelerates top- k queries for DNN interpretation. DeepEverest, with various optimizations, reduces the number of activations computed at query time with low storage overhead, while guaranteeing correct query results. With less than 20% of the storage of full materialization, DeepEverest accelerates individual queries by up to 63.5 \times and consistently outperforms other methods over various multi-query workloads.

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