STATISTICAL PATTERN RECOGNITION FINAL PROJECT REPORT

MAXIMUM LIKELIHOOD APPROXIMATE NEAREST NEIGHBOR METHOD IN REAL-TIME FACE RECOGNITION

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Abstract

Image recognition is one of the fundamental tasks in computer vision. Modern face recognition which is one of the substantial subtasks of the image recognition, in most cases, uses brute-force nearest neighbor method to iterate through the reference faces database to find the desired identity of the input image. Nowadays, most of these methods extract feature maps of the images using Convolutional Neural Networks. In this project, we'll be using Convolutional Neural Networks to extract image feature maps. We'll also analyze the implementation process of the Approximate Nearest Neighbor method to find the best ID for the input image.

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Introduction

1.1 Motivation

One of the well-known issues of vision systems building, is a processing of large databases. Unfortunately, nearest neighbor (NN) rule and exhaustive search usually cannot be implemented in real-time applications[1]. To overcome these problems, we'll analyze some of the purposed methods in this this topic.

1.2 Recent Methods of Image Recognition in Large Scale Databases

In this section we'll take a brief look at the recent researches on image recognition systems and their improvements for large scale databases.

1.2.1 Approximate Nearest Neighbor Shape Indexing

This method relies on the rapid recovery of the nearest neighbors from the index. In high-dimensional databases, standard k-d tree search often performs poorly. having to examine a large fraction of the points in the space to find the exact nearest neighbor. However, a variant of this search which efficiently finds approximate neighbors will be used to limit the search time. The algorithm, which we have called Best Bin First (BBF) search, finds the nearest neighbor for a large fraction of queries, and finds a very good neighbor the remaining times [2].

1.2.2 Directed Enumeration Method

Directed enumeration method is proposed to improve image recognition performance. The method is applied with similarity measures which do not met metric properties. This method increases performance in 3 to 12 times in comparison with nearest neighbor. Recognition speed using DEM is increased when many neighbors are located at similar distances [3].

1.2.3 Directed Enumeration Alternatives Modification

In this method, a new modification of the method of directed alternatives enumeration using the Kullback Leibler discrimination information is proposed for half-tone image recognition. Results of an experimental study in the problem of face images recognition with a large database are presented. It is shown that the proposed modification is characterized by increased speed of image recognition (5-10 times vs exhaustive search) [4].

Problem Definition and Algorithm Definition

In this chapter, we'll take a close look at the core algorithm of MLANN method. The next is dedicated to the implementation procedure of the algorithm.

2.0.1 MLANN; General Idea

Despite the most of known fast approximate NN algorithms, the proposed method is not heuristic. The joint probabilistic densities (likelihoods) of the distances to previously checked reference objects are estimated for each class. The next reference instance is selected from the class with the maximal likelihood. To deal with the quadratic memory requirement of this approach, the author has proposed its modification, which processes the distances from all instances to a small set of pivots chosen with the farthest-first traversal. Experimental study in face recognition with the histograms of oriented gradients and the deep neural network-based image features shows that the proposed method is much faster than the known approximate NN algorithms for medium databases [5].

2.0.2 How Statistical Face Recognition Works

In face recognition, we are required to assign an observed image X to one of R classes which is specified by the database of reference images. In this method, feature maps extracted from observed and reference images are treated as probability distributions. The following subsection provides baseline assumptions for this task.

Key Assumptions

In face recognition task with statistical method, we assume that

- 1. The reference images from different classes are independent;
- 2. The probability distributions of the feature vectors from the same class are identical [5].

2.0.3 Where the Problem Arises

The core algorithm of many face recognition implementations use Nearest Neighbor method as the main approach to find the most similarity between the input image and the reference images in the database. Let X be the observed image and X_r $r \in 1,...R$ be each of the images in the reference database. In that case, the optimal maximal likelihood solution of the face recognition task is achieved by

$$W_v: v = \arg\min_r \ \rho(X, X_r)$$
 (2.1)

where r is selected in a *brute-force* manner by the nearest neighbor algorithm and ρ is mean to be the *distance* of two image feature maps.

2.0.4 Speeding up the Search Process

Approximate Nearest Neighbor

To speed-up the search process, we can use *approximate* techniques. As an example, an approximate method is provided in [6]. This method is based on the following criterion:

$$W_v: \rho(X, X_v) < \rho_0 \tag{2.2}$$

which is the termination condition of the ANN method with respect to a bound of of ρ_0 .

Best-Bin First kd-tree

In this method, the first reference image X_{r_1} , $r_1 \in \{1, ..., R\}$ is randomly chosen. Next, it is put into the priority queue if reference images sorted by the distance to X. Next, the highest priority item X_i is pulled from the queue and the set of reference images $X_i^{(M)}$ is determined by using the following expression:

$$(\forall X_k \in X_i^{(M)})(\forall X_j \notin X_i^{(M)})|\rho_{i,k} - \rho(X, X_k)| \le |\rho_{i,j} - \rho(X, X_j)|$$
 (2.3)

the distance matrix $P = [\rho_{i,j}]$ is computed at the beginning of the algorithm.

Maximum Likelihood Approximate Nearest Neighbor Method

Here is the core algorithm of this paper. The method is proposed for the next step reference image choice. Let the reference images $X_{r_1}, ..., X_{r_k}$ have been checked before the k-th step, meaning that the distances $\rho(X, X_{r_1}), \rho(X, X_{r_2}), ... \rho(X, X_{r_k})$ have been computed. We can choose the next most probable reference image $X_{r_{k+1}}$ as

$$r_{k+1} = argmax(p_v. \prod_{i=1}^k f(\rho(X, X_{r_i})|W_v)) \quad v \in \{1, ...R\} - \{r_1, ...r_k\}$$
 (2.4)

where p_v is the prior probability of class v which is the class of the observed image. $f(\rho(X, X_{r_i})|W_v)$ is the conditional density of the distance $\rho(X, X_{r_i})$ if the hypothesis W_v is true. We can acquire the likelihood $f(\rho(X, X_{r_i})|W_v)$ using a Homogeneity Testing Probabilistic Neural Network. Using an small modification proposed in the paper, we can obtain

$$r_{k+1} = argmin(\sum_{i=1}^{k} \phi_{\mu}(r_i) - \ln p_{\mu}) \quad \mu \in \{1, ...R\} - \{r_1, ...r_k\}$$
 (2.5)

where $\phi_{\mu}(r_i)$ satisfies all we need for a normal distribution:

$$\phi_{\mu}(r_i) = \frac{(\rho(X, X_{r_i}) - \rho_{\mu, r_i} - \frac{N-1}{UV}))^2}{\rho_{\mu, r_i}} + \frac{4}{UV} \ln(2\rho_{\mu, r_i} + \frac{N-1}{UV})$$
(2.6)

since the average image size is usually much higher than the number of parameters UV >> (N-1), $\phi_{\mu}(r_i)$ can be simplified:

$$\phi_{\mu}(r_i) = \frac{(\rho(X, X_{r_i}) - \rho_{\mu, r_i}))^2}{\rho_{\mu, r_i}}$$
(2.7)

If the distance $\rho(X, X_{r_{k+1}})$ is lower than the threshold ρ_0 , then the search procedure is terminated at the $K_{checks} = k+1$ step. Otherwise, the reference image $X_{r_{k+1}}$ is put into the set of previously checked reference images and the maximal likelihood search procedure is repeated.

2.0.5 MLANN; Disadvantages

Though the proposed MLANN method makes it possible to minimize the average number of distance computations K_{checks} , it has two significant disadvantages:

- The necessity to compute and store distances between all reference images makes MLANN to use *quadratic memory space*.
- The complexity of extra computation at each step is rather high. For instance, $\phi_{\mu}(r_i)$ is calculated for each unchecked μ -th reference image.

2.0.6 Pivot Based MLANN; An Improvement to MLANN Method

In this section a modified version of MLANN is proposed. We'll choose $p \ll R$ reference images (pivots) from the database. The results show that the performance increases if we choose images far from each other. Thus it is necessary to choose the pivots as distant as possible. This task can be solved using known clustering algorithms. Another approach is the incremental selection of pivots which is better compared to the clustering techniques which requires calculating the distances between all the images in database. We'll use the farthest-first traversal procedure. The first pivot X_{r_1} is chosen randomly. Next, the distances between this reference image and all other reference images are computed, and the most distant reference image is selected as the second pivot. The procedure is repeated so that every pivot is characterized with the highest sum of distances to the previous pivots:

$$r_{i+1} = argmax \sum_{j=1}^{i} \rho(X_{\mu}, X_{r_j}), i \in \{1, ..., p-1\}$$
(2.8)

where

$$\mu \in \{1, ...R\} - \{r_1, ...r_i\}$$

The result of the initialization phase is a list of pivots and R * p matrix of distances between all reference images and the pivots.

2.0.7 Time and Space Complexity Analysis

Figure 2.1 demonstrates the time and space complexities of P-ML-ANN method.

	Time complexity	Space complexity
NN (exhaustive search)	$O(R \cdot N)$	$O(R \cdot N)$
Ordering permutations	$O(E_{max}\cdot(N+p)+plogp)$	$O(R \cdot (N+p))$
DEM	$O(E_{max}\cdot(N + Mlog(E_{max}\cdot M)))$	$O(R \cdot (N+R))$
ML-ANN	$O(E_{max}\cdot(N+R))$	$O(R \cdot (N+R))$
P-ML-ANN	$O(E_{max} \cdot N + Rlog E_{max})$	$O(R \cdot (N+p))$

Figure 2.1: Time and space complexities of the P-ML-ANN algorithm.

Implementation and Performance Analysis

3.1 Analysis of Naive Template Matching

In this section, we'll analyze the implementation and the performance of the *Naive Template Matching* algorithm. First of all, let's consider the serial implementation case.

3.1.1 Bitmap Images

For this project, we've used the *bitmap* image library by *Arash Partow* which is accessible here.

3.1.2 Serial Implementation of NTM

The serial implementation of this algorithm consists of 2 main loops to iterate over image pixels. There is also a need for each pixel we are iterating through, to calculate the Sum of Absolute Deviations which is the similarity measure between the main image pixels and the template image. Please refer to the Naive Template Matching algorithm to see the code. Since the task is mainly focused on counting the number of occurance inside the main images, we can obtain this by finding the minimum value of SAD while iterating through the main image pixels and counting the number of occurrences of the found minimum inside the main image matrix.

3.1.3 Timing the Serial NTM

We can obtain the elapsed time for the serial iteration using the *chrono* library. The code for this section is provided here. Please refer to the serial implementation code for the details.

3.1.4 CUDA Implementation of NTM

In this section we'll walk through the kernel codes provided in the kernel.cu.

Compute Sad Array Kernel

The image data is being stored in an 1-D array. Thus, the best choice for CUDA kernel launch parameters would be to set grid dimensions as $(image_width/block_size_x, image_height/block_size_y, 1)$. The main purpose of this choice is not the way we have stored the data, it actually depends on the input data. Since we are dealing with images it is more suitable to have 2-D blocks of threads. Further explanations focus on the kernel implementation. Initially, each thread finds its own global 2-D identification as rows and columns. Then, each thread virtually sees a kernel (template) around it self. This is the core of the parallelization section. All threads can see all of the iterations in one place. We then compute the SAD of each pixel and load it inside an SAD array. The next section describes the reason to do so.

Find Minimum in Array Kernel

We can obtain the best possible match by finding the minimum SAD in SAD array. We'll do so using shared memory to speed up the process. Each thread helps loading the data inside a block into the shared memory of that block. We then use *fminf* to find the minimum inside each block. Since we are using the shared memory, we need to do a *reduction* to find the minimum across different blocks. We can obtain this reduction task defining a *mutex* to control the global memory write by first thread of each block.

Find Number of Occurrences

Since the main task is to find the number of occurrences of template image in main image, we should count the number of occurrences of minimum SAD in the SAD array. To speed up the process we've used shared memory again. The intuition of using shared memory is exactly the same as the previous section.

3.1.5 Launch Parameters on Test Cases

The test has been done on three different sizes of images. One of the medium sized images is a bitmap image with dimensionality of 1112*1500. The template image is a 116*116 image in this case. Considering a block size of 1024 on a Nvidia~850m~GTX (compute capability of 5), the launch parameters are provided below:

	X	Y	Z
Grid Dimensions	38	47	1
Block Dimensions	32	32	1

3.1.6 Occupancy Analysis

Before getting into the details, it is important to mention that maximum number of active warps in theory is 64. *Nsight* profiler provides great details on the number of active warps per SM. In this case, there are 63.83 active warps on the first kernel (Compute Sad Array Kernel). The occupancy can obtained by dividing the number of active warps to the number of maximum warps available:

$$occupancy = \frac{63.83}{64} = 99.74\%$$
 (3.1)

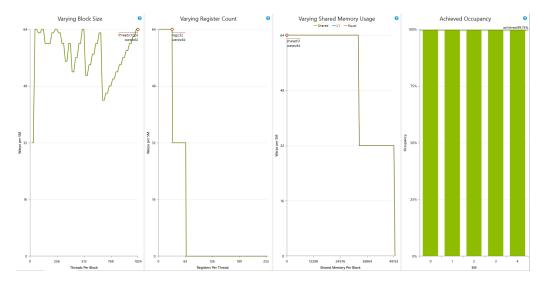


Figure 3.1: Illustration of varying the size of blocks, registers and shared memory on the achieved occupancy.

According to figure 3.1, varying the number of threads per block might yield the same result in other scenarios. The circled point shows the current number of threads per block and the current upper limit of active warps. Note that the number of active warps is not the number of warps per block (that is threads per block divided by warp size, rounded up). If the chart's line goes higher than the circle, changing the block size could increase occupancy without changing the other factors. Also, increasing the number of variables (registers per block) will drastically decrease the occupancy of the first kernel.

3.1.7 SM Activity Analysis

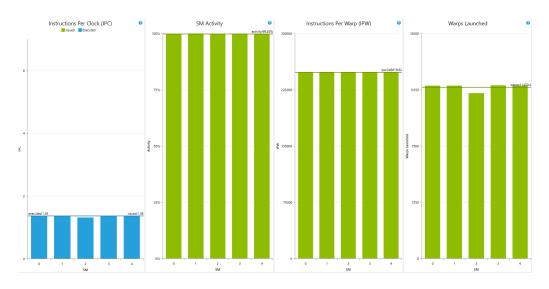


Figure 3.2: Issue efficiency and SM activity charts for the first (bottleneck) kernel.

The IPC chart demonstrates the achieved instructions throughputs per SM for both, issued instructions and executed instructions. The theoretical maximum peak IPC is a device limit and defined by the compute capabilities of the target device.

Issued IPC

The average number of issued instructions per cycle accounting for every iteration of instruction replays. Optimal if as close as possible to the Executed IPC. As described in the background section of this document, some assembly instructions require to be multi-issued. Hence, the instruction mix affects the definition of the optimal target for this metric. In this case, the issued

IPC throughput is minimum since there weren't any multi issue instructions in the first kernel.

Executed IPC

The average number of executed instructions per cycle. Each warp scheduler of a multiprocessor can execute instructions independently, a target goal of executing one instruction per cycle means executing on average with an IPC equal to the number of warp schedulers per SM. The maximum achievable target IPC for a kernel is dependent on the mixture of instructions executed.

SM Activity Chart

The second chart provides information on the activity of each SM during the kernel launch. A multiprocessor is considered to be active if at least one warp is currently assigned for execution. An SM can be inactive - even though the kernel grid is not yet completed - due to high workload imbalances. Such uneven balancing between the SMs can be caused by a few factors: Different execution times for the kernel blocks, variations between the number of scheduled blocks per SM, or a combination of the two. In this case, non of the above has occurred and the activity result is almost 100%.

IPW Chart

One of the most common patterns for varying IPWs is conditionally executed code blocks. IPW can be useful while having variations in SM activity chart which in this case, non of these have occurred.

3.1.8 Speedup Analysis

In this section, we'll analyze the possible theoretical speedup. For this purpose, we'll calculate the serial computation time for the *collection* and *collection_coin* images as described above. This table describes the average speedup of parallel naive template matching algorithm compared to serial implementation. As can be seen in the figure 3.3, the speedup obtained on GPU is dedicated to many different parameters. In order to calculate the theoretical speedup a direct use of Amdahl's law doesn't fit in this problem. One might say that the number of cores in this case is N (Same as the number of threads), but GPU does not have as many physical cores as the number of threads that can be launched. Additionally, significant portion of kernel time is begin spent just waiting for the data to be read/written from/to global memory. Also, the frequency and operational power of CPU

cores is higher than GPU cores. In this section, we'll use a modified version of the Amdahl's law [?]:

$$S = \frac{1}{(1-p) + (k*p/N)} \qquad k = freq(CPU)/freq(GPU)$$
 (3.2)

The core frequency of my Intel 4710HQ processor is 3490 MHz. The core frequency of each GPU core is 902 MHz. These information are extracted using beloved CPU-Z and GPU-Z programs. Replacing these information into 3.2 yields

$$S = \frac{1}{(1-p) + 3.86 * p/N} \tag{3.3}$$

since the size of input is constant in this law, we'll assume that 3.3 is correct for the third experiment. Let's suppose that $\frac{1}{4}$ of the main program is parallelized, thus $p=\frac{1}{4}$ and the number of threads to run is 1828864 (1786 blocks of 1024). Theoretical speedup will be

$$S = \frac{1}{\left(\frac{3}{4}\right) + 3.86 * \frac{\frac{1}{4}}{1828864}} = 1.33 \tag{3.4}$$

	Exp 1	Exp 2	Exp 3	Exp 4
Image * Template	(630 * 459) * (116 * 116)	(1203 * 460) * (116 * 116)	(1211 * 1500) * (116 * 116)	(3840 * 2160) * (214 * 214)
Serial NTM	10787 msec	22615 msec	92204 msec	1722131 msec
Parallel NTM	339 msec	644 msec	1975 msec	27983 msec
Parallel NTM + Data Transfer	345 msec	665 msec	1980 msec	28977 msec
Achieved Speedup	31.82	35.11	46.68	61.54

Figure 3.3: Computation and data transfer time for naive serial and parallel implementation.

3.1.9 Performance Analysis

In this section, we'll analyze the performance of the Compute Sad Array Kernel. Firstly, lets find the arithmetic intensity of this kernel. Arithmetic intensity I, also called operational intensity, is the ratio of arithmetic operations or work (W), to the memory traffic (Q) [?]:

$$I = \frac{W}{O} \tag{3.5}$$

and denotes the number of operations per byte of memory traffic. When the work is represented as FLOPS, the arithmetic intensity will be FLOPS/Byte.

According to the *Naive Roofline Model*, the *attainable performance* is bound either by the *peak performance* or *memory bandwidth* * *arithmetic intensity*. In this case, the operational intensity can be obtained by following the memory access patterns and the work being done in the main kernel: It is given

Statement	Operations	# Global Memory Writes * bytes	# Global Memory Reads* bytes	# Constant Memory Reads* bytes	# Shared Memory Reads* bytes
row, col	()()(*+*+)	0	0	0	6 * sizeof(int)
1st conditional	00(< <)0	0	0	2 * sizeof(int)	0
1st loop	00(<)0	0	0	1 * sizeof(int)	0
2nd loop	00(<)0	0	0	1 * sizeof(int)	0
m_r	()()(*)()+++*+	0	1 * sizeof(unsigned char)	1 * sizeof(int)	0
m_g	()()(*)()+++*+	0	1 * sizeof(unsigned char)	1 * sizeof(int)	0
m_b	()()(*)()+++*+	0	1 * sizeof(unsigned char)	1 * sizeof(int)	0
<i>t_r</i>	()()(*)() + * +	0	1 * sizeof(unsigned char)	1 * sizeof(int)	0
t_g	()()(*)() + * +	0	1 * sizeof(unsigned char)	1 * sizeof(int)	0
t_b	()()(*)() + * +	0	1 * sizeof(unsigned char)	1 * sizeof(int)	0
SAD Write conditional	()()(*<)() +	1 * sizeof(int)	0	1 * sizeof(int)	0
	5 / (6 * sizeof(unsigne	ed char) + 1 * sizeof(int) = 0.5			
Nvidia 850m GTX DDR Bus Frequency	1001 MHz				
Nvidia 850m GTX DDR Bus Bandwidth	2 * 128 = 256 bits				
Nvidia 850m GTX Memory Bandwidth	32 GB/s				

Figure 3.4: Calculation of first kernel operational intensity and GPU memory bandwidth.

that the *single precision* processing power of a *Nvidia 850m GTX* is 1155 GFLOPS [?]. Given these results, the kernel is actually memory bound since 0.5 < 1155/32 = 36. The kernel arithmetic intensity should be at least 36 to be considered a compute bound kernel.

3.2 Implementation of FFT Based Template Matching

In this section, we delve into the details of *FFT* based template matching using *cufft* library from *Nvidia*. Note that this implementation is still in progress and there are multiple serial loops conducted in preprocessing step which slows down the process compared to the *naive template matching* procedure. As mentioned before, the overall procedure is to convert the spatial domain of input images to the frequency domain. In the frequency domain, we can apply a correlation based operation. This operation yields a signal of the same size as main image size. We then apply an inverse Fourier transform to turn the results back to the spatial domain. We then find the maximum of that signal and the number of occurrences of that maximum value in that

signal. The procedure is well defined below[?]:

```
c = real(IFFT_{2D}(FFT_{2D}(main\_image).*FFT_{2D}(template\_image))) (3.6)
```

according to 3.6, c contains a list of values which we are desired to find the maximum of it (for maximum similarity). To obtain this in CUDA, we've used multiple functions which are described further. Note that * symbol shows the complex conjugate of the second signal.

3.2.1 Real to Complex Conversion

For this part, we have not focused on the performance. We use simple for loops to convert the image data matrix to the proper format of cufft library which is called cufftComplex. cufftComplex is a simple typedef of float type in C.

3.2.2 Padding Data

Before performing a 2-D Forward Fourier Transform we are urged to perform a data padding on template signal to make it the same size as the main signal. Figure 2.1 illustrates this phenomenon pretty well. The duty of data padding is simple. Allocate new memory with a new size. Copy signals into their corresponding location in new allocated memory and set other places to 0.

3.2.3 2-D FFT Planning

cufft library uses the concept of plan to provide the baseline setup for the main APIs provided in the library. A plan can initiate a 1-D, 2-D or 3-D data processing API from cufft library. In this implementation, we need a setup for a 2 dimensional Fourier transform, thus we use cufftPlan2d to prepare the library for the main transformations.

3.2.4 Forward FFT Launch

In order to launch a forward *complex to complex* Fourier transform using *cufft*, we'll use *cufftExecC2C*. The constant *CUFFT_FORWARD* shows the direction of the Fourier transform which is *Forward* in this case.

3.2.5 Complex Conjugate Kernel

This kernel implements the conjugation procedure. The implementation is pretty straightforward. Each thread multiples the corresponding signal point's complex part to -1 and loads it back to the input signal.

3.2.6 Complex Point-wise Kernel

This kernel provides a point to point multiplication manner for each of the points in main and template input signals. It uses another kernel called *ComplexMul* to perform a *complex multiplication*. It then scales the result using *ComplexScale*. Scaling or normalization is mainly done by dividing the signal values to the size of the signal.

3.2.7 Inverse FFT Launch

Finally, we'll perform an inverse Fourier transform on the result of the *Complex Point-wise Kernel*. The resulting signal is in the spatial domain.

3.2.8 Maximum Value and Number of Occurrences

To find the number of occurrences, we *serially* iterate over the resulting signal of the previous section. Note that these steps are not 100% efficient and are still in improvement.

Future Work

We admit the possible drawbacks and inefficiencies existing in this project, thus we have the following key improvement to-do list:

- Reorganization of the project structure, header files and .c files.
- Reorganization of the project source code for a C unified implementation.
- \bullet Usage shared memory techniques to improve the performance of the NTM implementation.
- Usage of highly-optimized image libraries such as *OpenCV* for *image* rotation, *image* matrix extraction and general bitmap processing.
- Implementation of CUDA kernels for real-to-complex signal conversions.
- Implementation of CUDA kernel to find the number of occurrences of template signal in main signal in Fourier based template matching.

Conclusion

In this project, we analyzed various algorithms for the task of *Image Template Matching*. It is fair to say that there are already highly optimized libraries with template matching capabilities such as OpenCV, but as mentioned before, the main purpose of this project is to improve the understanding of the CUDA environment and parallel implementation of highly-demanded computer vision tasks such as template matching. We analyzed two main methods to perform this task:

- 1. Naive Template Matching
- 2. FFT Based Template Matching

For further read, please refer to methods such as Stereo Matching, Image Registration and Scale Invariant Feature Transform.

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