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**Qualification work  
to obtain a Master's degree**

in the educational and scientific program "Artificial Intelligence"  
specialty 122 "Computer Science" on the topic:

**ASSOCIATIVE METAMEMORY**

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## ABSTRACT

The master's thesis consists of an introduction, three chapters, conclusions, a list of used sources (45 items), and 1 appendix. The work contains 15 figures and 5 tables. The total volume of the work is 32 pages, the main text of the work is laid out on 19 pages.

O(1) KNN, NEUROHASHING, INDEXES, VAN EMBDE BOARS TREE, INFORMATION RETRIEVAL, ASSOCIATIVE MEMORY, METAMEMORY, FAST DATA STRUCTURES, SOURCE KNOWLEDGE, RETRIEVAL-AUGMENTED, SELF-SUPERVISED.

The object of research is artificial episodic memory. The subject of research is associative metamemory.

The objective of the work is to develop an approach for quick searching and saving a multidimensional manifold of memories in a one-dimensional (linear) memory, which is inherent in all computers implementing the von Neumann architecture.

Development methods: multimodal artificial neural networks, similarity search methods for dense vectors, code quantization methods, proximity graph indexing methods. Development tools: Visual Studio and PyCharm development environments, C++ and Python 3 programming languages, VS Performance Profiler.

The results of the work: the necessity of using neurohashing has been proven, along with the inability of conventional algorithmic indexing methods to find and add a large number of vector representations in real-time as the database grows in size, a model has been trained, which returns a binary number that forms a linear order; a data structure has been implemented for fast neighbor search.

## ABBREVIATIONS

$k$ -NN	–	$K$ -nearest neighbors algorithm
Faiss	–	Facebook AI Similarity Search
Glass	–	Graph Library for Approximate Nearest Search
GPU	–	Graphics Processing Unit
HNSW	–	Hierarchical Navigable Small World
IVF	–	Inverted File Index
LLaMA	–	Large Language Model
LSH	–	Locality-Sensitive Hashing
mAP	–	Mean Average Precision
MIPS	–	Maximum Inner Product Search
NGT	–	Neighborhood Graph and Tree
NMSLIB	–	Non-Metric Space Library
OPQ	–	Optimized Product Quantization
ScaNN	–	Scalable Nearest Neighbors

## INTRODUCTION

### Assessment of the current state of the development object

Transformers [1] is the leading architecture of artificial neural networks for modeling and predicting sequences. In the simplest case, generative transformers learn to predict the next token using the information of all previous tokens within the window. For example, words embedded in a multidimensional vector space or small rectangular fragments of an image ( $n, m$ ) can serve as tokens. They are decomposed into one-dimensional vectors ( $n * m$ ) and inserted into the vector space according to the principle similar to words [2].

One of the approaches to improving the output quality of generative transformers is the use of an external knowledge base (retrieval-augmented) from which initial knowledge is taken (source knowledge). In the case of text transformers, relevant sentences are taken from the database. Relevance is determined by the scalar product (MIPS), cosine similarity, or Euclidean distance between nested vectors of these sentences. REALM is a prominent representative of this approach [3], DPR [4], RAG [5], FID [6], RETRO [7], R2-D2 [8], and Atlas [9]. These architectures mainly solve the problem of answering open questions.

In general,  $k$  nearest neighbors ( $k$ -NN) are searched for the relevant elements retrieval. Episodic memory is a cognitive analog of the knowledge base, metamemory is an analog of  $k$ -NN. Metamemory plays one of the key roles in thinking: a person can immediately, without much thought, say that he does not know the fifth largest dinosaur. This allows you to significantly save brain resources because otherwise, it would be necessary to recall everything that a person knows to conclude that he does not know something.

In addition to knowledge, nested vectors from previous contexts can be stored in the database. This is done in Memorizing Transformers [10] and Unlimiformer [11] architectures. This implicitly expands the current context to a size limited only by the quality of the search.

## **Relevance of the work and the reasons for its implementation**

The number of vectors in the knowledge base can reach billions. For fast search, modern architectures use multi-level similarity indices based on quantization and graph search. Their query time increases linearly as  $k$  increases. Empirically, it allows us to find the order of thousands sentence 64 tokens each without significant impact on training time. This amount of text is not enough for some tasks, for example, summarizing (referencing). In the field of computer vision, this is even more noticeable, because a thousand images is only ~17 seconds of video at a frequency of 60 frames per second.

Apparently, the human brain does not spend a lot of time on the assimilation of sensory information. In other words, indexing happens instantaneously as new information is integrated. Mind-wandering also does not require a long search. Instead, commonly used  $k$ -NN algorithms are sublinear at best, such as LSH [12] –  $O(n^{p < 1})$ .

## **Objective and tasks of the work**

The objective of the work is to develop an approach for adding and searching the knowledge base in constant time, using neurohashing and fast data structures. To achieve this goal, the following tasks have been set:

- measure the running time of the most used similarity indices on synthetic data
- analyze available neurohashing methods
- adapt one of them for multimodal mode
- compare popular datasets
- index one of the datasets
- select a fast data structure for this index

## SECTION 1. SIMILARITY INDEXES

### 1.1. Comparison of similarity search methods

From search libraries based on peer-reviewed papers, the most famous are Meta Faiss [13] and Google ScaNN [14], from non-academic – Spotify Annoy, NMSLIB, Kakao N2, Yahoo NGT. In addition to open libraries, there are many semi-commercial vector databases. Among them: Milvus > Weaviate > Qdrant (ranked by speed) and Pinecone (not tested).

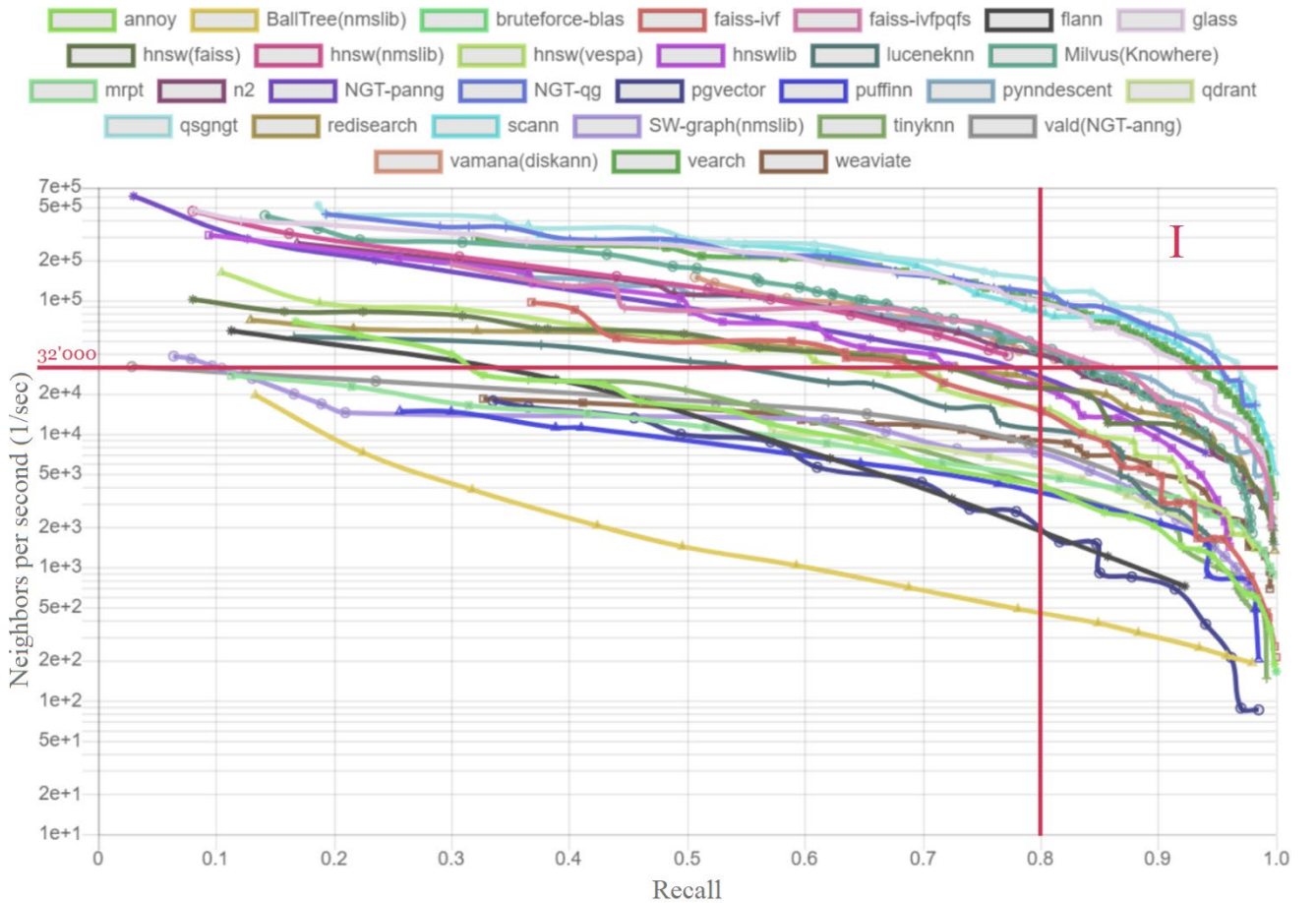


Figure 1 – Performance measurements of a large number of methods<sup>1</sup>.

Red lines demarcate the 1st quadrant: number of queries > 32'000, recall > 0.8

A priori, we will set the lower performance limit of 32,000 requests per second with a recall of at least 0.8. For small  $k$ , this amount is sufficient for real-time execution. Then, after filtering, 11 methods will remain:

<sup>1</sup> <https://github.com/erikbern/ann-benchmarks/>  
<https://ann-benchmarks.com/>

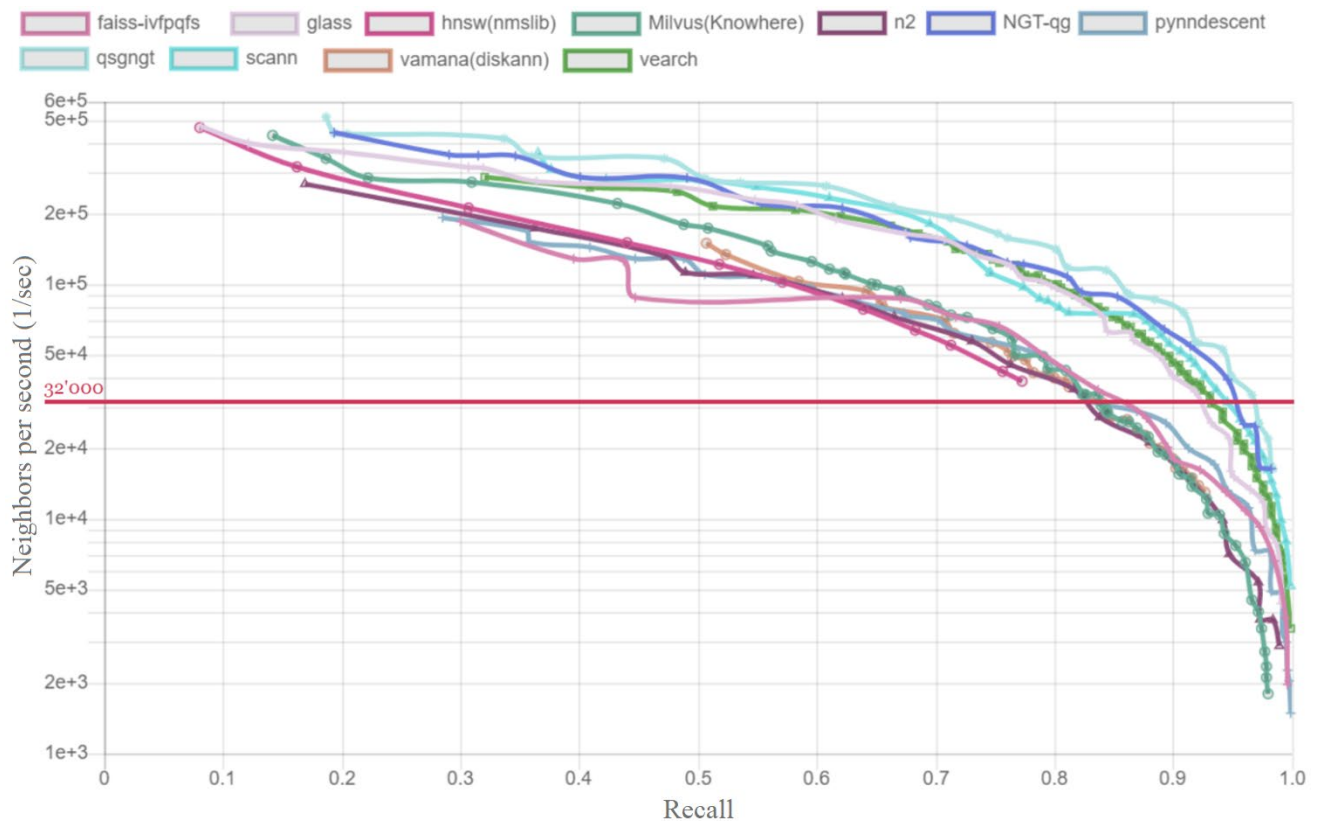


Figure 2 – Performance measurements of real-time methods

The list of real-time methods includes the previously reviewed Faiss and ScaNN. A group of the fastest methods clearly stands out on the graph: qsgngt > NGT > Glass > ScaNN. Among them, NGT and ScaNN are widely used.

## 1.2. Description of the Faiss library

Faiss indices are less memory-intensive than ScaNN, because ScaNN, in addition to storing the quantized index, needs to store the original vectors. Also, Faiss has better documentation. Therefore, it is advisable to start research with Faiss.

Description of Faiss from the repository: "Faiss is a library for efficient similarity search and clustering of dense vectors. It contains algorithms that search in sets of vectors of any size, up to ones that possibly do not fit in RAM. It also contains supporting code for evaluation and parameter tuning. Faiss is written in C++ with complete wrappers for Python/numpy. Some of the most useful algorithms are implemented on the GPU. It is developed primarily at Meta's Fundamental AI Research group".



One of the main methods for index formation is quantization. It is used to efficiently find similarities in high-dimensional data. The goal of quantization is to reduce the dimensionality of a dataset while preserving important information about the data. This provides faster and more efficient search algorithms that can be used to speed up tasks such as image and text searches.

In Faiss, quantization is performed by dividing the dataset into several smaller subspaces [15], which are called Voronoi cells. Each Voronoi cell corresponds to a cluster of similar data points. Next, the centroids of these clusters are calculated and stored as a lookup table. When a query is performed on a dataset, it is first quantized into one of the Voronoi cells and then finds the nearest centroid.

The main advantage of quantization is that it reduces the amount of data that needs to be stored and searched, making it much faster than other methods. In addition, quantization can be easily parallelized, making it suitable for large-scale data analysis.

Faiss supports the comparison of Euclidean distances, scalar products, and cosine similarities. All three metrics are proportional to each other:

$$\begin{aligned}\cos(\theta_{xy}) &= \frac{\langle x, y \rangle}{\|x\| \|y\|} \\ \|x - y\|_2^2 &= \sum_{i=1}^n (x_i - y_i)^2 = \|x\|_2^2 + \|y\|_2^2 - 2x^T y = 2(1 - \cos(\theta_{xy})) \\ \langle x, y \rangle &\propto \cos(\theta_{xy}) \propto \|x - y\|_2^2\end{aligned}$$

Euclidean distance should be used when the norms of the vectors make sense.

### 1.3. Faiss index factory

The Index Factory is a helper module in Faiss that provides a convenient way to create and configure index objects. The factory takes care of index initialization, training, and persistence, and provides a simple interface to configure various index parameters, such as the number of clusters, the number of bytes per code in quantization, and the number of levels.

In this work, the following configuration was used for testing:  
*OPQ16\_64,IVF1000\_HNSW32,PQ16x4fs*

OPQ [16] aims to reduce the distortion introduced by vector quantization by applying an orthogonal transformation to the input vectors before quantization. The transformation rotates and scales the vectors in such a way as to minimize the distance between the transformed vectors and the centroids of the clusters, resulting in more accurate quantization.

The OPQ algorithm works by first partitioning the input vectors into multiple blocks and then iteratively optimizing the rotation and scaling of each block to minimize the quantization error. Optimization is performed using gradient descent, with gradients calculated from an analytical expression.

*OPQ16* at the beginning of the configuration indicates only the rotation matrix, so at the end, we must have *PQ16x4fs*, which is responsible for the quantization process [15] and means 16 codes of 4 bits in *fast mode scan*. Thus, the code size is  $16 * 4 / 8 = 8$  bytes. The *\_64* in *OPQ16\_64* is an additional dimensionality reduction operation to 64. *IVF1000\_HNSW32* indicates a combination of 1000-cell IVF and HNSW [17], where each centroid is connected to 32 others.

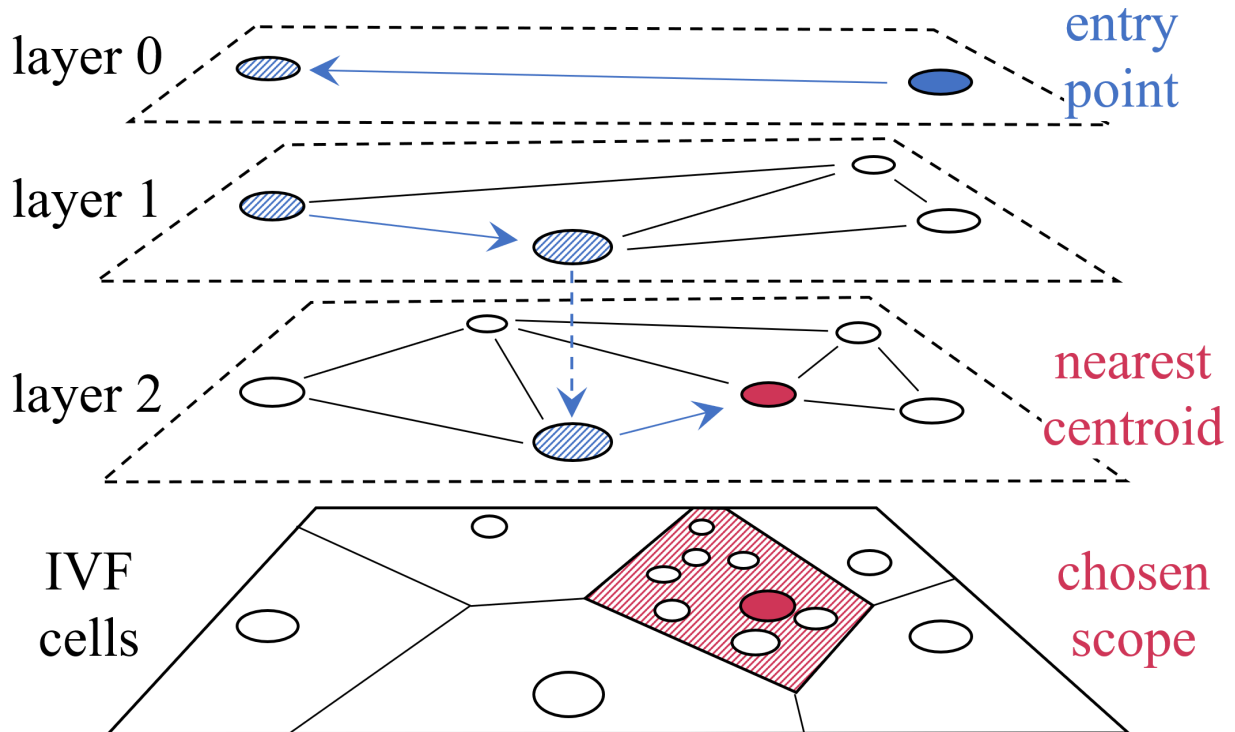


Figure 3 – HNSW and IVF. A replica from the Pinecone blog

### 1.4. Testing of indexes

To guarantee obtaining  $k$  neighbors in IVF, it is necessary to make samples on neighboring cells because the method is approximate. Moreover, with increasing  $k$  the number of required samples is also increasing. In Faiss, the `nprobe` parameter is responsible for the number of samples.

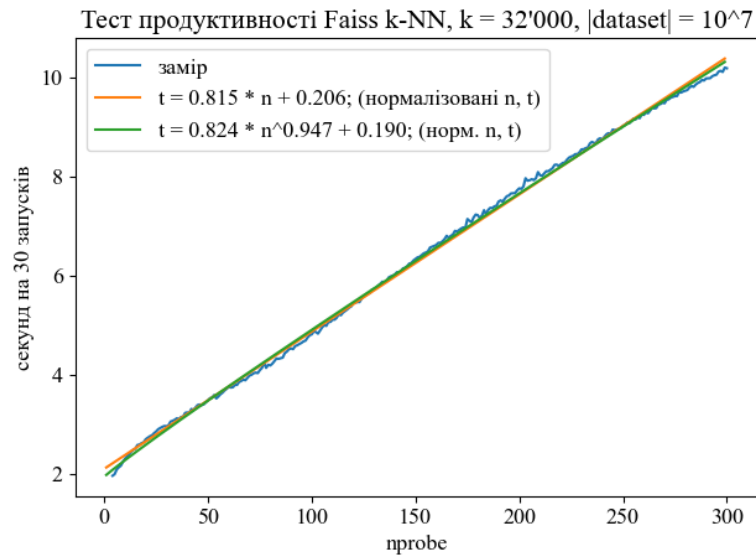


Figure 4 – Dependence of query time on `nprobe`

– In the legend of the graph, approximations of the measurement by polynomials of the first order and power functions with a shift are given. As you can see, the query time increases linearly (we will consider the nature of the power of 0.947 to be false).

The next step is to check the dependence of the request time on  $k$ :

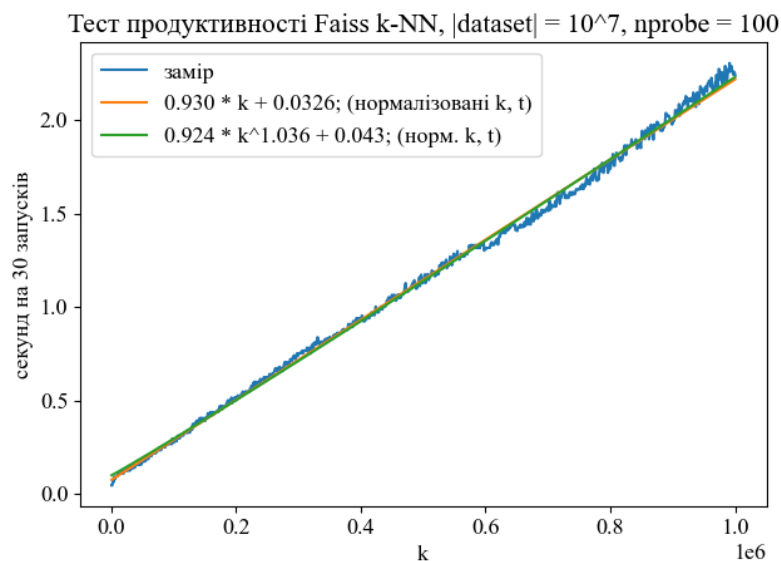


Figure 5 – Dependence of the request time on the number of neighbors in  $k$ -NN

This dependence also turned out to be close to linear (we will consider the nature of the power of 1.036 to be erroneous). Instead, the dependence on the size of the database goes from sublinear to superlinear as  $k$  increases ( $0.639 \rightarrow 0.755 \rightarrow 0.911 \rightarrow 1.01 \rightarrow 1.14$ ):

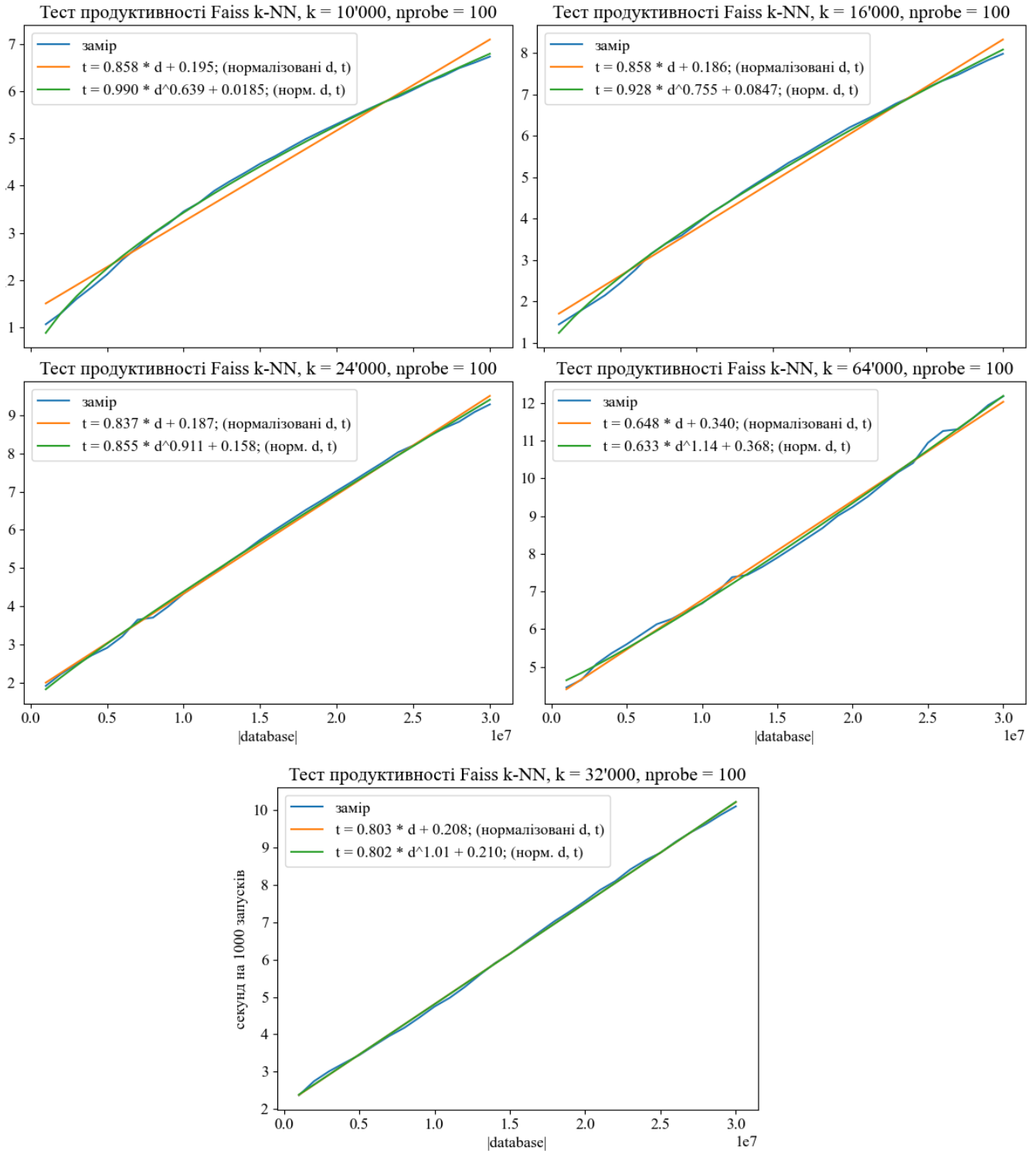


Figure 6 – Dependence of query time on database size

Given the recent advances in increasing the size of the context to  $\sim 2$  million tokens [18], let's calculate how long the query will take in the case of the text dataset The Pile [19], which after processing contains 5.8 billion fragments of 64 tokens each. This is equivalent to  $k$ -NN with  $k = 2'048'000 / 64 = 32'000$ .

That is, we can use the non-normalized analog  $t = 0.803 * d + 0.208$ :

$t = 2.7 \text{ e-}07 * 5.8\text{e}9 + 2.1 \approx 1568 \text{ seconds}/1000 \text{ starts} = 1.568 \text{ seconds/request}$ . This is exactly on schedule faiss-ivfpqfs in Fig. 2 and measured completeness 0.89. From the same graph, it can be seen that the fastest methods ScaNN and NGT-qg are ahead of Faiss by up to three times.

This duration significantly slows down training. In the article that introduces the LLaMA generative transformer [20], the authors noted that the throughput of the model for 65 billion parameters is 380 tokens/second/GPU on 2048 A100 GPUs with 80 GB of video memory. That is, the bandwidth of the cluster is 778'240 tokens/second. If we go back to the  $\sim 2\text{M}$  long context, that's 2.634 seconds/request.

Besides slowing down training by  $1.568 / (1.568 + 2.634) = 37\%$  in the case of Faiss and 17% in the faster methods, the similarity indices have quite a few counterintuitive hyperparameters. At the same time, artificial neural networks are quite often ahead of classical algorithms. Their direct propagation has a time constant within a few milliseconds. This suggests that there should be a neural network approach that works in  $k * O(\text{const}) = O(k)$ .

## SECTION 2. NEUROHASHING

### 2.1. Intuition

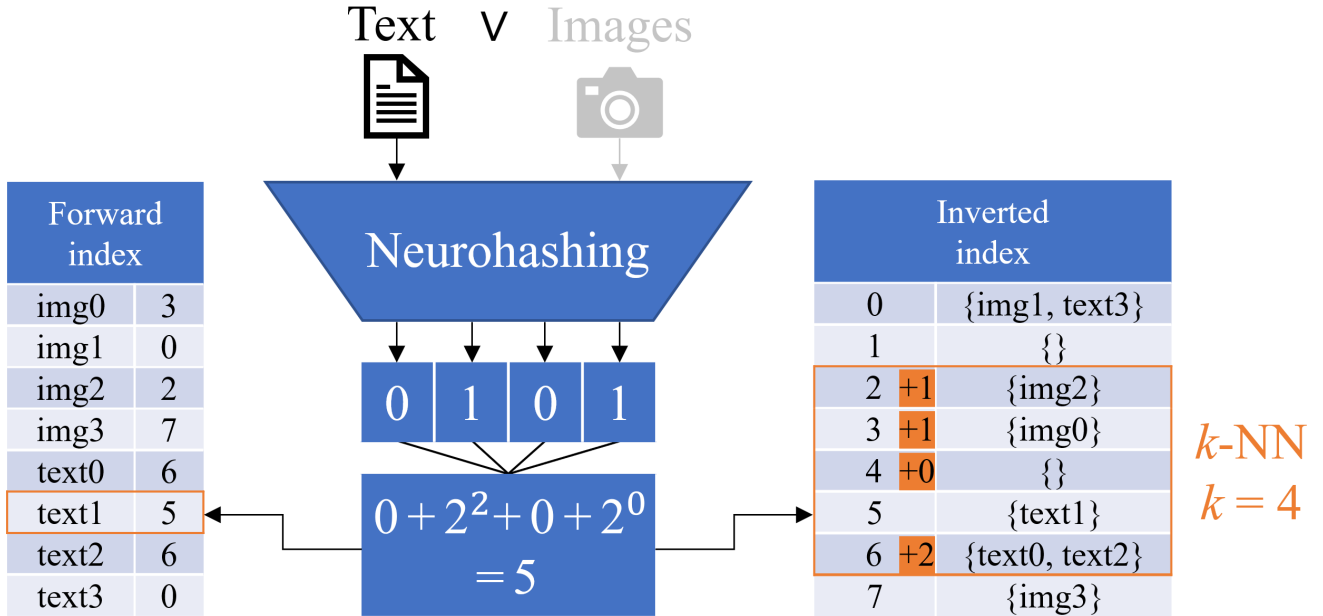


Figure 7 – Concept of architecture. "text1" is added to the direct and inverted index under number 5. For  $k$ -NN, it remains to greedily go through the inv. index

The idea is to model a hierarchy of universals (features) using a hierarchy of numbers. So that the binary number 0100 (4) is semantically closer to 0110 (5) than to 1100 (12). To show that this is possible, consider the case of one-dimensional  $k$ -NN:

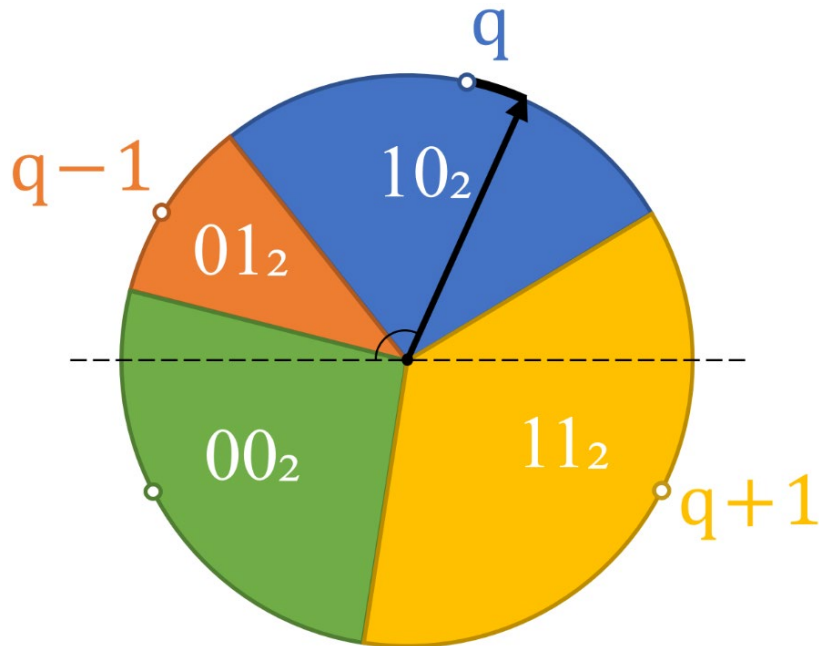


Figure 8 – Two-dimensional projection of a one-dimensional Voronoi diagram

Colored zones are responsible for belonging to a certain vertex. For example, the black query vector has entered the cell of the blue vertex. A neural network can easily learn the dependence of the index (color) on the angle of the vector. Binary network outputs are a natural way to learn a binary numerical index. If there are many points, and the numbers of the cells are arranged according to the time arrow, then it is not difficult to search for  $k$  nearest neighbors by index – you need to take vertices with indices from  $q - k / 2$  to  $q + k / 2$ , where  $q$  is the index of the query vector.

In the two-dimensional case, everything is much more complicated, because the ground truth order of vertices by distance is discontinuous, and therefore we have a decision boundary with self-intersections. This complicates learning and impairs the generalization abilities of the neural network. This can be compensated to some extent by having a large, balanced dataset during pretraining because such a set will force the neural network to efficiently compress the representation.

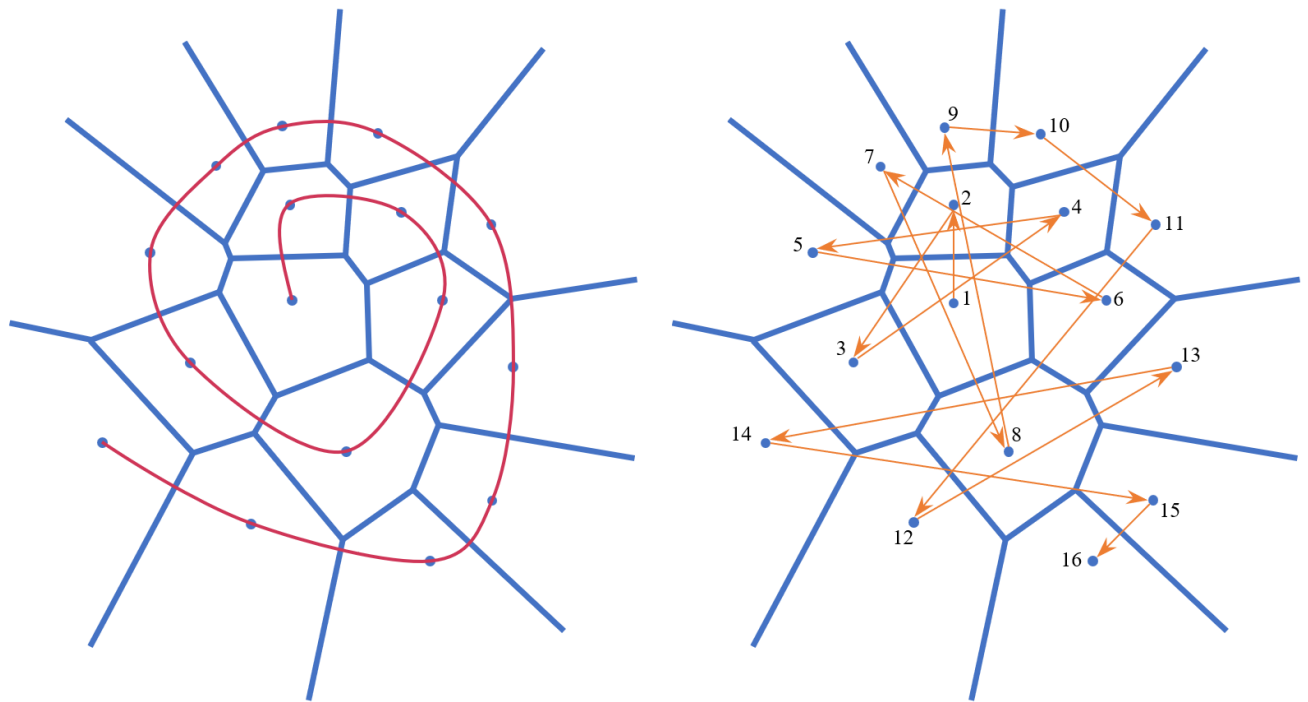


Figure 9 – Two-dimensional Voronoi diagram.

On the left is the desired smooth order of the vertices by distance from the first one, on the right is the discontinuous true order. A neural network learns the true order

## 2.2. Data description

Most works use three sets of images: CIFAR-10 [21], NUS-WIDE [22], MS COCO [23]. There are various filtered subsets and partitions of these sets. In CIFAR-10 each image has one of 10 classes, in NUS-WIDE-21 each image has up to 21 classes, in MS-COCO-2014 up to 80 classes. The set is divided into three parts: a training part (train), a test part (query), and a database (database). During testing a binary code (hash) is generated for the request image, which is searched in the database. Thus, the database can contain training images, because they do not overlap with test images.

The balancing of samples underwent changes during the development of the direction. Recent works by TBH [24], NSH [25], WCH [26] converged on the same principle: CIFAR-10 – sampling with uniform distribution, NUS-WIDE-10 – each class has at least 100 images, MS-COCO-2014 – unbalanced sampling. Some past work either neglected to balance the dataset or contained errors. For example, in HashNet [27] the sample database contains query, which may result in perfect results on certain elements. Therefore, the hashing code was taken from the public DeepHash repository<sup>2</sup> after being checked for correctness.

Architectures; Dataset	Query (Test)	Train	Database	Total
HashNet; NUS-WIDE-81-m	5'000 (unbalanced)	10'000 (unbalanced)	Train + Database0 = 218 ' 491	223'496
HashNet; MS-COCO-2014	5'000 (unbalanced)	10'000 (unbalanced)	<b>Test!</b> + Database0 = 112 ' 218	122'218
CIBHash; CIFAR-10	5'000 (not included) (+ 5'000 valid)	500 * 10	Train + Database0 = 50 ' 000	60'000
TBH, NSH, WCH; CIFAR-10	1'000 * 10	5 ' 000 * 10	Train = Database = 50 ' 000	60'000
TBH, CIBHash, NSH, WCH; NUS-WIDE-21	2100 (min. 100/class)	10'500 (min. 100/class)	Train + Database0 = 193 ' 734	195'834
TBH, CIBHash, NSH, WCH; MS-COCO-2014	5'000 (unbalanced)	10'000 (unbalanced)	Train + Database0 = 117 ' 218	122'218

Table 1 – Comparison of samples

<sup>2</sup> <https://github.com/swuxyj/DeepHash-pytorch>



### 2.3. CIBHash Binary Siamese Neurohashing

In these datasets, we have true labels (image classes), but we will use them only for testing. Because we aim to train a model without supervision (control). Such methods of machine learning are called self-supervised. One of the popular self-supervised approaches to studying vector representations is the use of dual architectures. They are also called Siamese architectures, distillation approaches, joint embedding architectures, and methods of mutual information maximization.

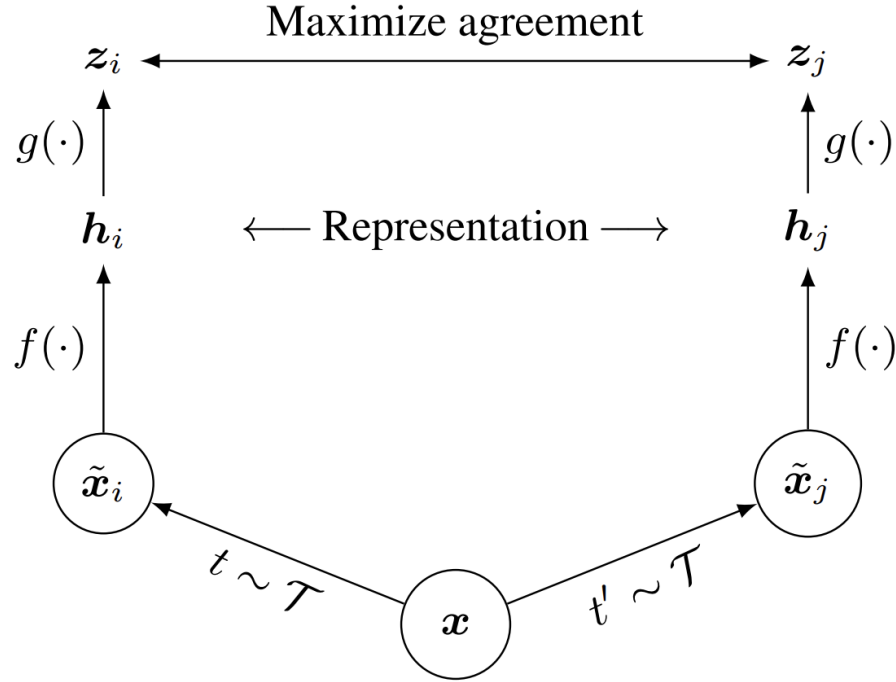


Figure 10 [28] – Contrast learning architecture SimCLR.

$t$  and  $t'$  these augmentations are obtained from one family  $\mathcal{T}$ ;  $f(\cdot)$  is a coding network,  $g(\cdot)$  is a projection head used only during training

Most of them use contrast learning to avoid collapsing to a trivial solution. However, this requires a large selection of negative elements [29]. Instead, there are methods that do not require a negative sample. Instead, they use different regularizations to prevent collapse. We will present the ranking of these methods by classification accuracy on ImageNet-1k [30]:

DINOv2 [31] > EsViT [32] = I-JEPA [33] > DINO [34] >= VICReg [35] > Whitening-based methods [36] = Barlow Twins [37] > BYOL [38]

Moreover, the methods that are oriented to the modality of the dataset (in the case of ImageNet, this is an image) show themselves to be more accurate: DINOv2.. DINO. It makes no sense to adapt them to the text because their architecture is based on specific augmentations of images and other aspects not inherent in texts. VICReg..BYOL multimodal methods are less precise but more versatile. Practice shows that over time too specific methods give way to universal ones<sup>3</sup>, but such methods apparently do not yet exist.

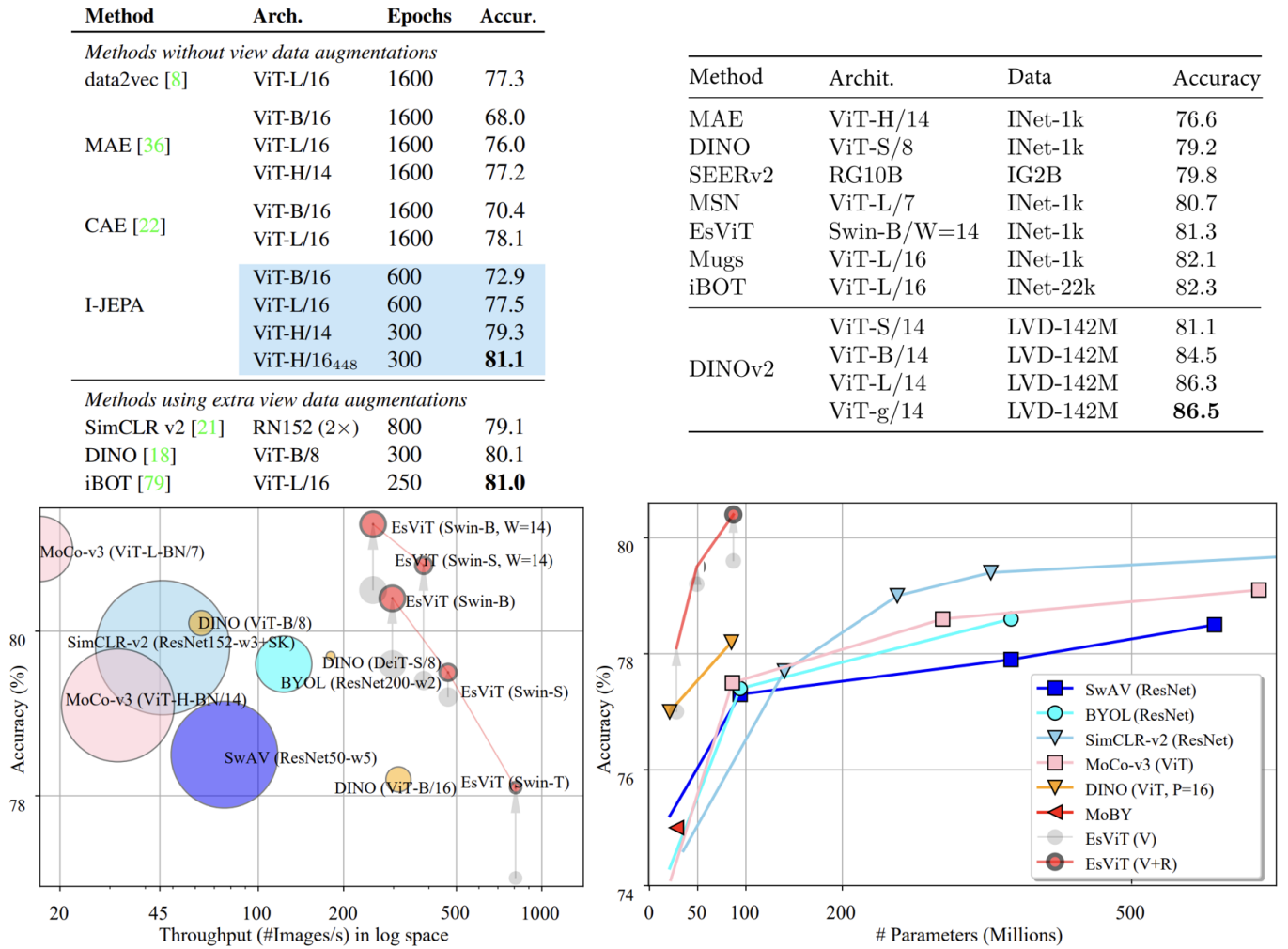


Figure 11 – Confirmation of ranking. Extract from articles [33], [31], [32]

Unfortunately, we cannot use the listed dual self-supervised methods, because their nested vectors consist of floating-point numbers. Comparing such vectors is too slow. Therefore, neurohashing methods use binary codes. Conventional artificial

<sup>3</sup> <http://www.incompleteideas.net/IncIdeas/BitterLesson.html>

neural networks, by their very nature, are not capable of returning discrete values. However, there are many mathematical approaches that allow us to bypass this limitation. A comparison of binary neurohashing methods is provided in the review [39]. Among the latest, it is worth mentioning CIBHash [40], NSH [25], WCH [26], and SDC [41]. The further these methods depart from the original SimCLR architecture [28], the more there is an unjustified tendency to unnatural statistical approaches and various mathematical tricks.

In this work, we will use CIBHash, because NSH does not have an open-source code, in WCH they train the basic feature extraction model, which is unfair, and SDC has too complex configurations. CIBHash is a fairly simple architecture: a basic overtrained VGG feature extraction model [42] which is frozen during training and has two linear layers. Binarization is achieved using a straight-through gradient estimator: when backpropagating the *sign* error, the function is simply skipped.

To achieve linear order, an error function was added converting from a binary code to a decimal number. That is, both parts of the dual model return one number, then these numbers are subtracted for the positive and negative elements of the sample. Positive elements are fragments of the same image and should have a smaller difference. Despite the simplicity of the approach, it should work well, because such an operation passes the gradient well.

The original feature selection model, or rather nested vectors, was replaced by the BEiT-3 multimodal model [43]. Potentially if text augmentations are added to CIBHash this will allow text to be indexed in addition to images. Also, BEiT performs better on image classification than VGG, and therefore its nested vectors contain more information about the input data. This should increase CIBHash's performance.

✓	✗	✗	✓	✓	mAP @5 = 1/2 * (AP <sub>1</sub> + AP <sub>2</sub> ) ≈ 1.62 = 0.81
P@1 = 1/1	P@2 = 1/2	P@3 = 1/3	P@4 = 2/4	P@5 = 3/5	
rel@1 = 1	rel@2 = 0	rel@3 = 0	rel@4 = 1	rel@5 = 1	
P@1 * rel@1 = 1/1	P@2 * rel@2 = 0/2	P@3 * rel@3 = 0/3	P@4 * rel@4 = 2/4	P@5 * rel@5 = 3/5	
AP <sub>1</sub> @5 = 1/3 * (1/1 + 0/2 + 0/3 + 2/4 + 3/5) = 0.7					
✓	✓	✗	✓	✗	
P@1 = 1/1	P@2 = 2/2	P@3 = 2/3	P@4 = 3/4	P@5 = 3/5	
rel@1 = 1	rel@2 = 1	rel@3 = 0	rel@4 = 1	rel@5 = 0	
P@1 * rel@1 = 1/1	P@2 * rel@2 = 2/2	P@3 * rel@3 = 0/3	P@4 * rel@4 = 3/4	P@5 * rel@5 = 0/5	
AP <sub>2</sub> @5 = 1/3 * (1/1 + 2/2 + 0/3 + 3/4 + 0/5) ≈ 0.92					

Table 2 – Example of  $\text{mAP}_{@5}$  calculation

Метод	CIFAR-10			NUS-WIDE			MS COCO		
	16 біт	32 біти	64 біти	16 біт	32 біти	64 біти	16 біт	32 біти	64 біти
AGH	0.333	0.357	0.358	0.592	0.615	0.616	0.596	0.625	0.631
ITQ	0.305	0.325	0.349	0.627	0.645	0.664	0.598	0.624	0.648
DGH	0.335	0.353	0.361	0.572	0.607	0.627	0.613	0.631	0.638
SGH	0.435	0.437	0.433	0.593	0.590	0.607	0.594	0.610	0.618
BGAN	0.525	0.531	0.562	0.684	0.714	0.730	0.645	0.682	0.707
BinGAN	0.476	0.512	0.520	0.654	0.709	0.713	0.651	0.673	0.696
GreedyHash	0.448	0.473	0.501	0.633	0.691	0.731	0.582	0.668	0.710
HashGAN	0.447	0.463	0.481	-	-	-	-	-	-
DVB	0.403	0.422	0.446	0.604	0.632	0.665	0.570	0.629	0.623
DistillHash	0.284	0.285	0.288	0.667	0.675	0.677	-	-	-
TBH	0.532	0.573	0.578	0.717	0.725	0.735	0.706	0.735	0.722
MLS <sup>3</sup> RDUH	0.369	0.394	0.412	0.713	0.727	0.750	0.607	0.622	0.641
DATE	0.577	0.629	0.647	0.793	0.809	0.815	-	-	-
MBE	0.561	0.576	0.595	0.651	0.663	0.673	-	-	-
CIMON	0.451	0.472	0.494	-	-	-	-	-	-
CIBHash	0.590	0.622	0.641	0.790	0.807	0.815	0.737	0.760	0.775
SPQ	0.768	0.793	0.812	0.766	0.774	0.785	-	-	-
NSH	0.706	0.733	0.756	0.758	0.811	0.824	0.746	0.774	0.783

Table 3 [26] – CIFAR-10:  $\text{mAP}_{@1000}$ ; NUS-WIDE, MS COCO:  $\text{mAP}_{@5000}$

## 2.4. BEiT multimodal transformer

Machine learning researchers have recently focused on developing basic, general-purpose models that can work across multiple modalities and adapt to a variety of downstream tasks. The Microsoft research group presented BEiT-3 – a multimodal base model for visual and speech-visual tasks. BEiT-3 interprets visual tokens as text in a foreign language that the authors call "Imglish" and performs unified masked "language" modeling on images, texts, and image-text pairs. The model learns on monomodal and multimodal data through a joint Multiway network Transformer, which uses a common self-attention module and different forward propagation networks for each modality. BEiT-3 achieved superior performance in various tests:

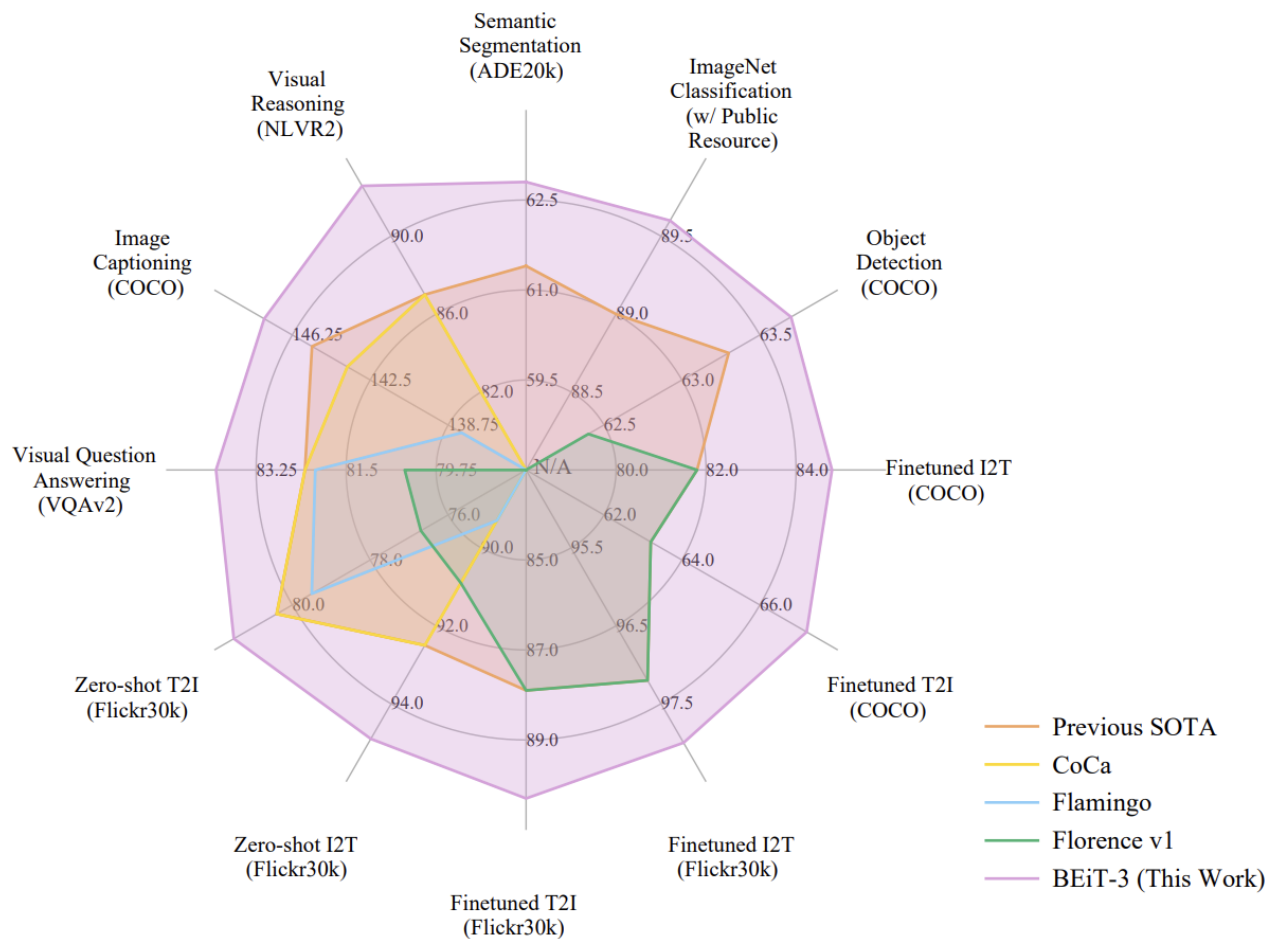


Figure 12 [43] – BEiT-3 is ahead of other basic models.

I2T/T2I = image-to-text/text-to-image search

## 2.5. Results of training

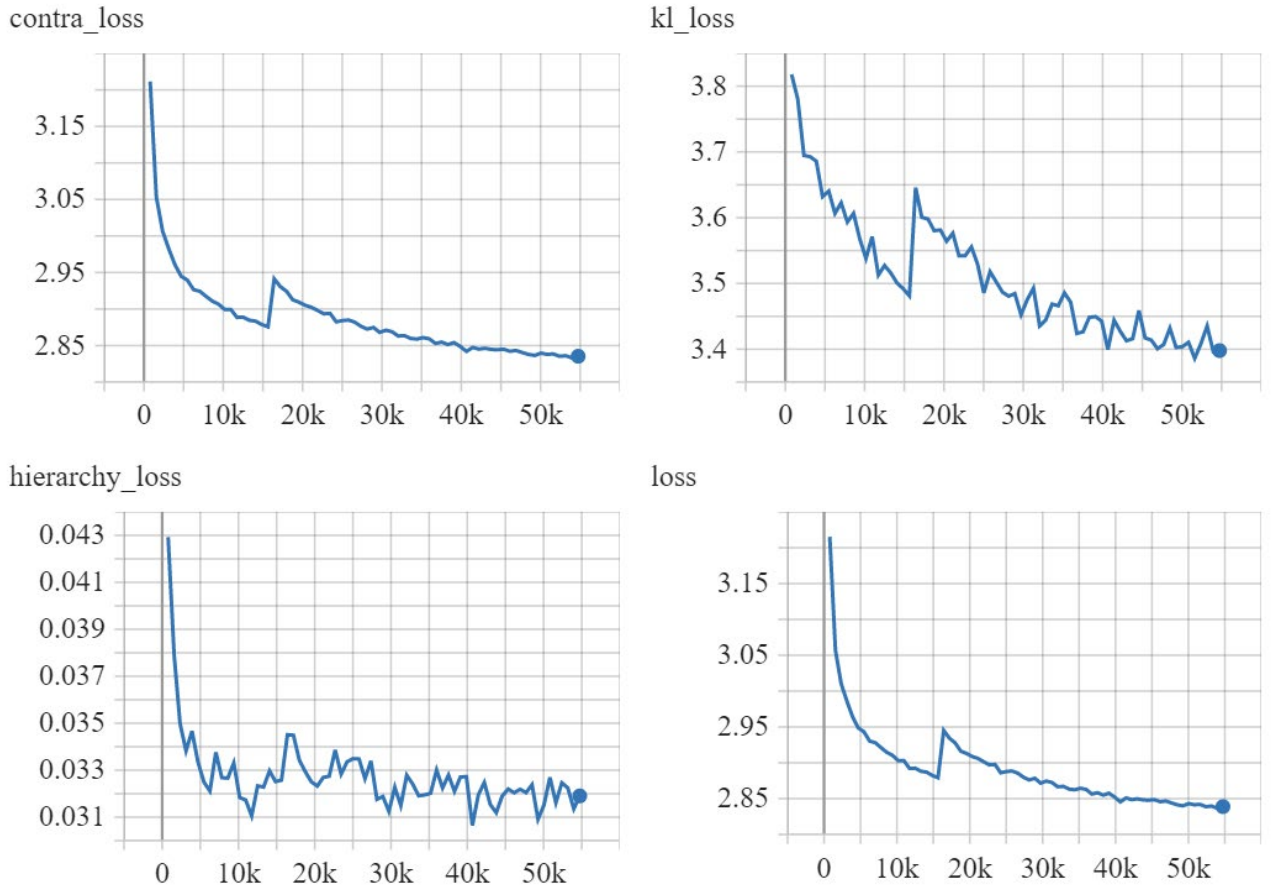


Figure 13 – Learning curves. Contra and KL loss functions were introduced in CIBHash, Hierarchy is the difference of decimal numbers introduced in this work.

The shift around 15k is caused by the restart

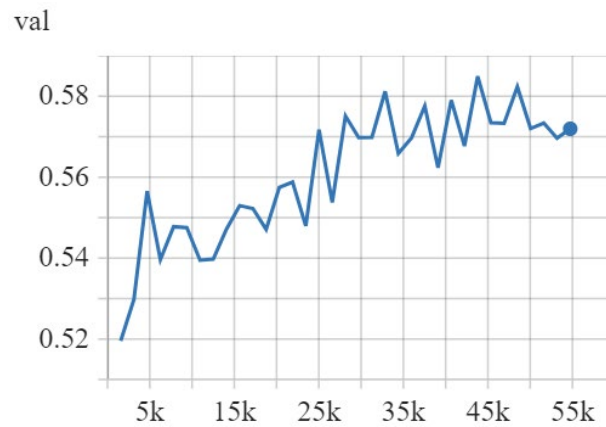


Figure 14 – Validation curve: mAP @1000

## SECTION 3. VAN EMDE BOARS TREES

### 3.1. Description of vEB trees

Let's say we can get a number for each item in the dataset. There is no guarantee that these numbers will not be the same for different items or guarantees that at least one item will match each number. That is, we have a large Boolean mask  $\sim$  a sparse array. Searching for neighbors in it is not as easy as in a completely filled array, because the greedy search can take  $O(n)$ . To avoid this, you can use van Emde Boas trees [44]. They are capable of doing addition and neighbor search in  $O(\log(\log(u)))$ , where  $u$  (universe) is the maximum number of elements. In our case, the length of the code returned by neurohashing is constant, and therefore the addition and search operations in the tree are also constant. Moreover, the constant is very small.

Unfortunately, decreasing time complexity inevitably leads to increasing memory complexity. An analogy can be drawn between these trees and counting sorting. In cases where speed is in the first place, memory is not paid attention to.

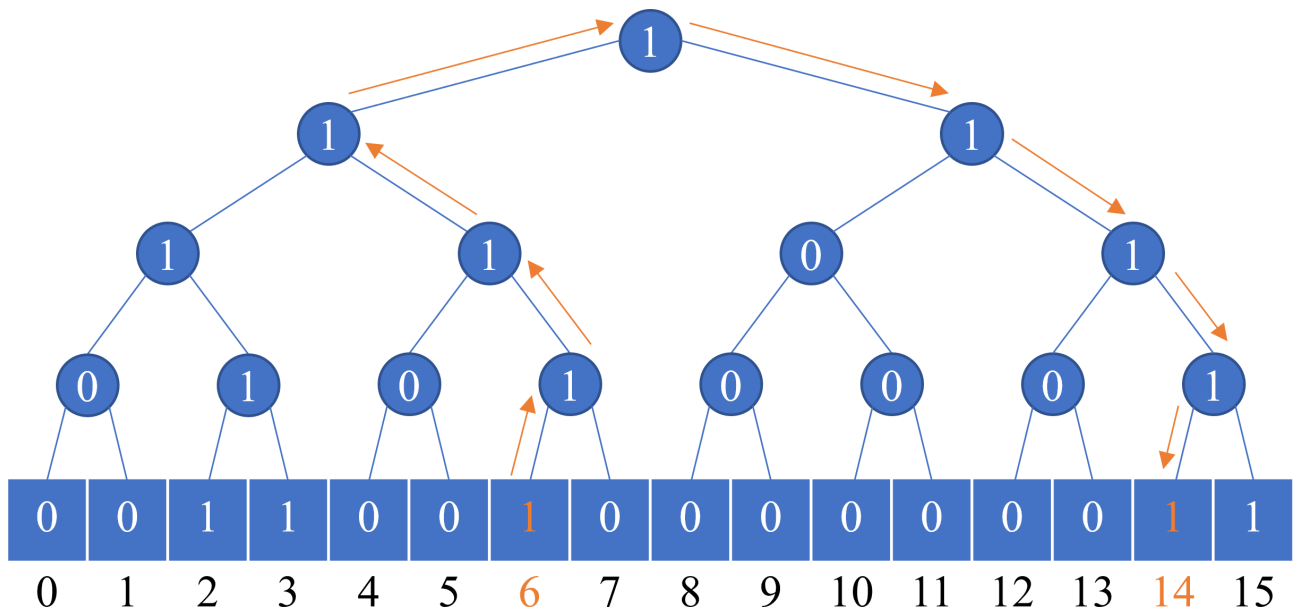


Figure 15 – Prototype of the vEB tree containing elements 2, 3, 6, 14, 15. The search for the successor of 6 is highlighted in orange. It is element 14

In fact, trees are more complex, because they have more than two branches at each vertex to save memory. They are very difficult to implement because operations on them have a non-trivial recursive nature. Also, vEB trees are not parallelized since the search for the next neighbor depends on the result of the previous search.

To reduce the search and append time constant, it is advisable to supplement the vEB tree with a doubly linked list and the following mapping (C++ map, Python dict): `{element_number: pointer_to_list_node}`.

Finally, we have such a course of action when adding:

1. We find the element number  $N$  using neurohashing
2. Using the tree, we set the predecessor and successor number  $N$
3. With the help of mapping, we find predecessor and successor nodes of  $N$  in the list
4. We replace them with the corresponding pointers to node  $N$

When searching, the procedure is the same, but in step 4, you need to run two greedy searches for  $k/2$  neighbors for the previous and next node of the list.

### 3.2. Testing of vEB trees

As mentioned earlier, trees are difficult to implement. Their memory and speed performance varies from implementation to implementation. To understand the limits of their performance, tests of various implementations found on the Internet were conducted. Some of them (opengenius, github/Petar) turned out to be inoperable. github/Julian is written in Python, all others in C++. Externally, github/Julian has very high-quality code and test coverage, but its results are worse than any of the C++ implementations.

The speed of addition and search has intervals because it depends on the number of elements in the tree. Two quantities were taken for testing: the minimum possible 32'000 and the maximum possible  $2^{24}$ . The test code is given in Appendix A. The obtained results confirm that with the help of vEB trees it is possible to query a large number of neighbors in real time.



Implementation	Memory used depth=24, GB	Addition speed k=32'000, 1/sec ~ Hz	Search speed k =32'000, 1/sec
opengen <sup>4</sup>	—		
github/Petar <sup>5</sup>	—		
github/Julian <sup>6</sup>	27.3	[1; 1]	[2; 11]
github/TISparta <sup>7</sup>	3.7	[38; 48]	[39; 96]
github/dragoun <sup>8</sup>	1.3	[15; 24]	[26; 49]
geeksforgeeks <sup>9</sup>	3.0	[34; 77]	[43; 85]

Table 4 – Measurements of different vEB trees. In terms of memory, the dragoun implementation is optimal, in terms of speed and memory – geeksforgeeks

Method	Element size	Addition k= 32'000, sec	Query k= 32'000, sec	Memory usage  db =5.8e9, GB
Naive: argsort(norm(database – query))	768 f32	1'254	10'535	17'817
argsort(hamming_dist(database – query))	32 b8	40	345	185.6
argsort(abs(decimal_database – d_query))	1 i32	4.7	740	23.2
Python boolean mask	1 b8	1/350	18.6	5.8
Python vEB tree (github/Julian)	—	1.09	0.35	6'978
C++ boolean mask	1 b8	1/1'000	[2.6e-4; 4]	5.8
C++ vEB tree (github/dragoun) + Mapping, doubly linked list + BEiT CIBHash	—	1/13 + 3.2e-7 + 3.84e-3 = 1/12	1/15 + 4.3e-4 + 3.84e-3 = 1/14	332.8 + 562.6 = 895.4
Faiss GPU (OPQ16_64,IVF100 0HNS...)	8 i8	1.25	1.57	88.7

Table 5 – Measurements of different  $k$ -NN approaches.

Unacceptable indicators are marked in red, mediocre ones in orange.

768 f32 – vector [768] of 32 bits float, 32 b8 – vector [32] of 8 bits boolean etc.

The transition between Faiss and the proposed doubly-linked list approach is interesting. An increase in speed by ~10 times leads to an increase in memory costs by ~10 times. It may seem that ~900 GB of RAM is an unaffordable amount, but at the moment Google has it Cloud has virtual machines up to 11776 GB. In our case, a VM with 976 GB costs from \$6 per hour<sup>10</sup>.

<sup>4</sup> <https://iq.opengenus.org/van-emde-boas-tree>

<sup>5</sup> <https://github.com/PetarV-/Algorithms>

<sup>6</sup> <https://github.com/Julian/veb>

<sup>7</sup> <https://github.com/TISparta/Van-Emde-Boas-tree>

<sup>8</sup> <https://github.com/dragoun/veb-tree>

<sup>9</sup> <https://geeksforgeeks.org/proto-van-emde-boas-tree-set-6-query-successor-and-predecessor>

<sup>10</sup> [https://cloud.google.com/compute/vm-instance-pricing#m3\\_machine\\_types](https://cloud.google.com/compute/vm-instance-pricing#m3_machine_types)

## CONCLUSIONS

A neural network approach was proposed that bypasses the existing methods of finding neighbors in speed. This can reduce the output time for a highly loaded question-and-answer system or enable real-time visual memory to be used by the agent during reinforcement learning. Because the best search libraries spend at least half a second for one query at  $k=32'000$  on The Pile. But this dataset is far from the largest available. If visual data will continuously flow with the frequency of the human eye from the virtual environment during reinforcement learning, then the speed of none of the libraries will be sufficient to search in such a volume of data.

Despite the high speed, the search accuracy remains quite low: 0.59 against Faiss' 0.89. Most likely, this is due to the fact that all neurohashing is reduced to training two linear layers. It makes sense to increase the number of layers and replace them with Gated MLP [45] because such layers show results similar to the attention mechanism while remaining simple in structure and fast in execution.

The image-specific neurohashing method WCH shows comparable accuracy to Faiss. This suggests that multimodal methods also have such potential.

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## APPENDIX A. TREE TESTING CODE

```

#include <cmath>
#include <iostream>
#include <vector>
#include <numeric>
#include <chrono>
#include <assert.h>
#include <cstdint>
#include <random>

#include "vEBTree.hpp"
#include "util.hpp"

using namespace std;

int main() {
    const int32_t sz = 1 << 24;
    const int32_t filled = sz;
    const int32_t nk = 32000;
    const int32_t log_every_nk = 500;

    assert(nk <= filled);
    assert(filled <= sz);

    vEBTree tree(sz);

    int32_t* v = new int32_t[sz];
    iota(v, v + sz, 0);
    auto rng = default_random_engine{};
    shuffle(v, v + sz, rng);

    auto a = chrono::high_resolution_clock::now();
    for (int32_t i = 0; i < filled; ++i) {
        tree.insert(v[i]);
    }
    auto b = chrono::high_resolution_clock::now() - a;

    float seconds = chrono::duration_cast<chrono::milliseconds>(b).count() / 1000.;
    cout << float(filled) / nk / seconds << endl;

    int32_t query, succ;
    a = chrono::high_resolution_clock::now();
    for (uint32_t i = 0; true; ++i) {
        query = v[rand() % (filled - 1)];
        succ = tree.successor(query);

        if (i % (log_every_nk * nk) == 0 && i) {
            b = chrono::high_resolution_clock::now() - a;

            if (filled == sz) {
                assert(succ == query + 1);
            }

            seconds = chrono::duration_cast<chrono::milliseconds>(b).count() / 1000.;
            cout << log_every_nk / seconds << ' ' << query << ' ' << succ << '\n';
            a = chrono::high_resolution_clock::now();
        }
    }

    return 0;
}

```