Carnegie Mellon University

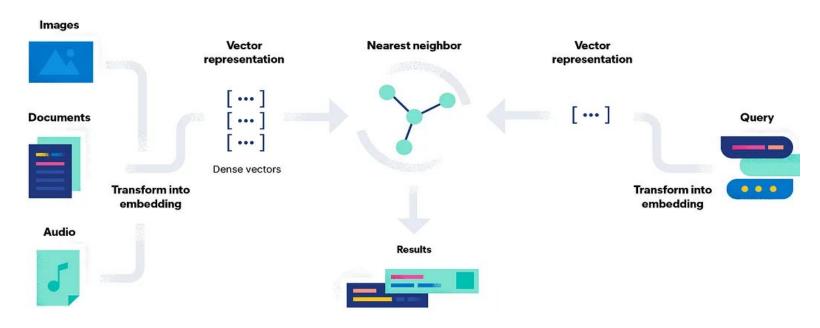
Efficient and robust approximate nearest neighbor search using Hierarchical Navigable Small World (HNSW) graphs

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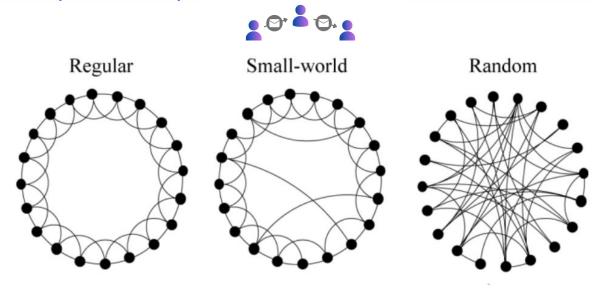
Motivation

- Similarity Search: applications in ML, retrieval, and with genAl -> RAG.
- KNN -> ANN: computational complexity vs. search accuracy.



Motivation for Navigable Small Worlds (NSW)

Six degrees of separation experiments run by Milgram in the 1960s.



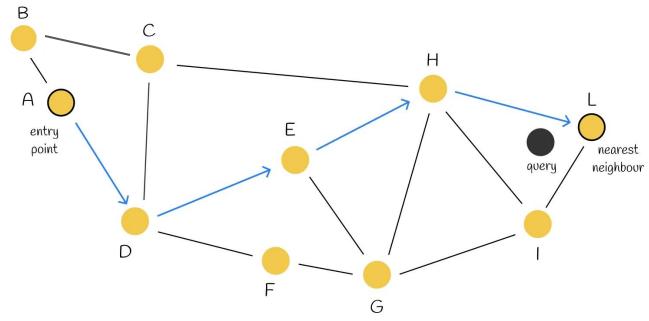
High clustering coefficient High distance

High clustering coefficient **Low** distance

Low clustering coefficient Low distance

ANN algorithm: Navigable Small Worlds (NSW)

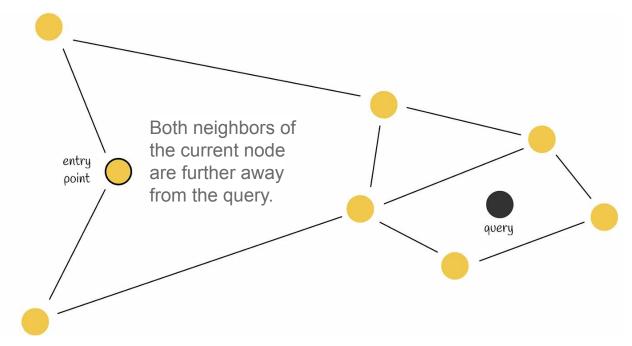
• O(log^k n) search and insertion, more useful for high dimensional large dataset



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ANN algorithm: Navigable Small Worlds (NSW)

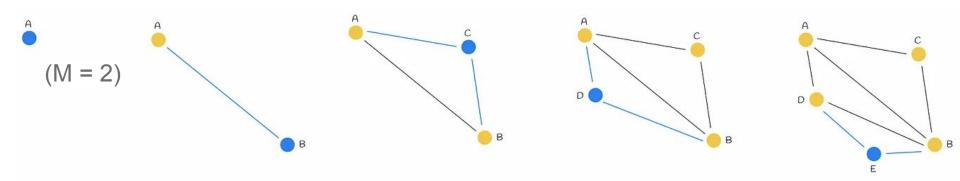
Greedy search can be trapped in local optimum (early stopping)



NSW Graph Construction

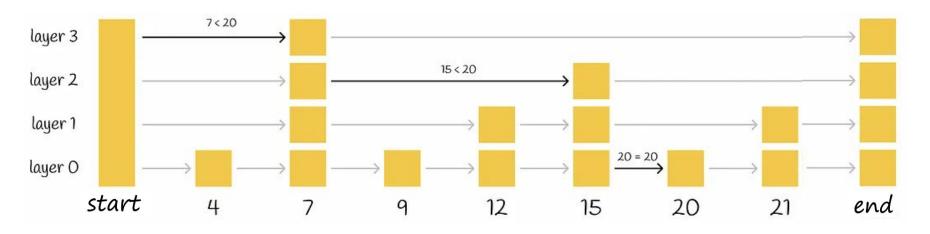
- Insert random points and link edges to M nearest neighbors (search)
- Longer edges are likely created at the beginning phase of graph construction

- "later become bridges between the network hubs that keep the overall graph connectivity and allow the logarithmic scaling of the number of hops during greedy routing."



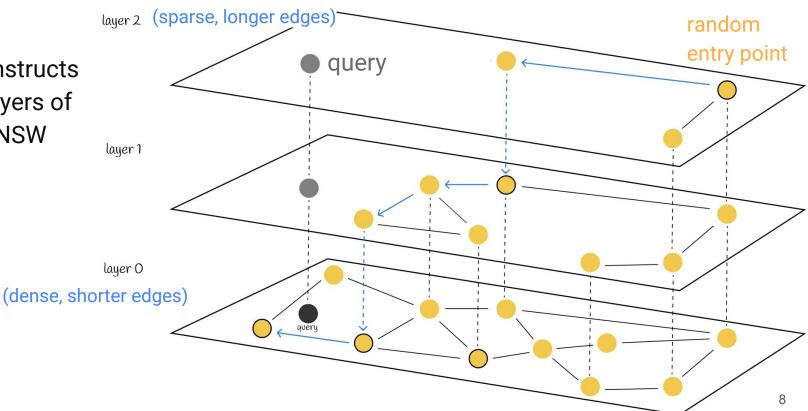
Data structure inspiration: Skip Lists

- O(log n) time complexity on average for both insertion and search
- Layered format with longer edges in the highest layers (for fast search) and shorter edges in the lower layers (for accurate search).

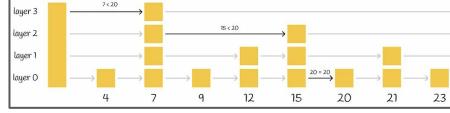


HNSW: Hierarchical Navigable Small Worlds

HNSW constructs multiple layers of proximity NSW graphs.



HNSW: Search



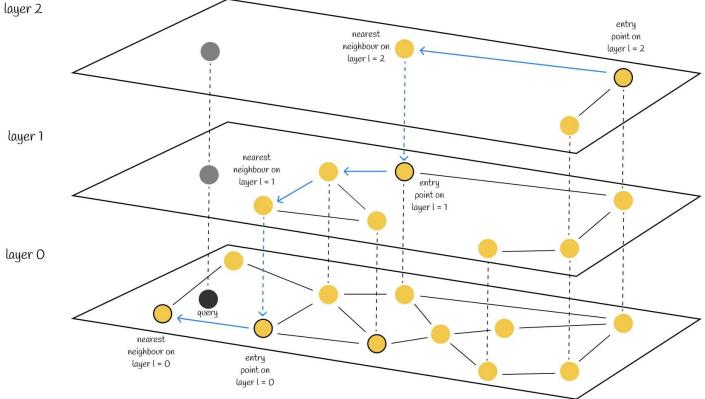


Figure: https://towardsdatascience.com/similarity-search-part-4-hierarchical-navigable-small-world-hnsw-2aad4fe87d37

HNSW: Search

- Based on principles of skip list and NSW
- Starts from the highest layer
- Proceeds to one level below each time, to find the local nearest neighbor among that layer nodes
- Return the nearest neighbor found on the lowest layer

Algorithm 5

K-NN-SEARCH(hnsw, q, K, ef)

Input: multilayer graph *hnsw*, query element *q*, number of nearest neighbors to return *K*, size of the dynamic candidate list *ef*

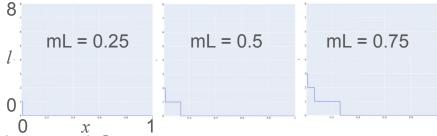
Output: *K* nearest elements to *q*

- 1 $W \leftarrow \emptyset$ // set for the current nearest elements
- 2 $ep \leftarrow$ get enter point for hnsw
- 3 *L* ← level of *ep* // top layer for *hnsw*
- 4 for $l_c \leftarrow L \dots 1$
- $W \leftarrow SEARCH-LAYER(q, ep, ef=1, l_c)$
- 6 $ep \leftarrow \text{get nearest element from } W \text{ to } q$
- 7 $W \leftarrow SEARCH-LAYER(q, ep, ef, l_c = 0)$
- 8 **return** *K* nearest elements from *W* to *q*

HNSW: Search

```
Algorithm 2
SEARCH-LAYER(q, ep, ef, lc)
Input: query element q, enter points ep, number of nearest to q ele-
ments to return ef, layer number lc
Output: ef closest neighbors to q
1 v \leftarrow ep // set of visited elements
2 C \leftarrow ep // set of candidates
3 W \leftarrow ep // dynamic list of found nearest neighbors
4 while |C| > 0
  c \leftarrow extract nearest element from C to q
  f \leftarrow \text{get furthest element from } W \text{ to } q
   if distance(c, q) > distance(f, q)
      break // all elements in W are evaluated
   for each e \in neighbourhood(c) at layer l_c // update C and W
10
     if e ∉ v
    v \leftarrow v \cup e
    f \leftarrow \text{get furthest element from } W \text{ to } q
     if distance(e, q) < distance(f, q) or |W| < ef
     C \leftarrow C \cup e
    W \leftarrow W \cup e
16 if |W| > ef
     remove furthest element from W to q
18 return W
```

Graph Construction



Q: How many layers can a node present in the graph?

A: Denote the maximum layer where the node can present as *l*

$$I = float[-ln(uniform(0, 1)) \cdot mL]$$

The number of layers *l* for every node is chosen randomly with exponentially decaying probability distribution

"To achieve the optimum performance advantage of the controllable hierarchy, the overlap between neighbors on different layers has to be small."

mL value tradeoff:

- a smaller mL: more traversals on each layer
- a larger mL: more overlaps

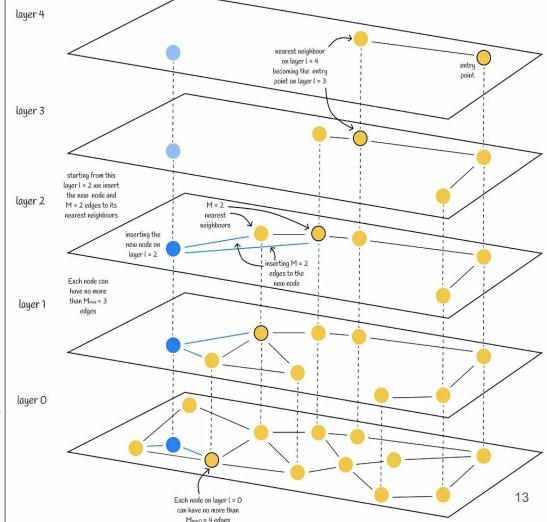
HNSW: Insertion

First phase:

- Greedily search for the nearest node from the upper layer
- 2. Use it as an entry point to the next layer until reaching layer *l*

Second phase:

- 1. Insert a new node from layer *l*
- Greedily search for *efConstruction* nearest neighbors
- 3. Choose M out of them and build edges
- 4. Use each of found *efConstruction* nodes as entry points to the next layer until layer 0



HNSW: Insertion

M value tradeoff:

A smaller M is better for lower recalls or low-dimensional data; A larger M is better for high recalls or high-dimensional data

efConstruction value tradeoff:

A larger value implies a more profound search as more candidates are explored, but requires more computations

 M_{max} the maximum number of edges a vertex can have

Algorithm 1

INSERT(hnsw, q, M, Mmax, efConstruction, ml) **Input**: multilayer graph *hnsw*, new element *q*, number of established

connections M, maximum number of connections for each element per layer Mmax, size of the dynamic candidate list efConstruction, normalization factor for level generation mu Output: update hnsw inserting element q

1 $W \leftarrow \emptyset$ // list for the currently found nearest elements

2 $ep \leftarrow$ get enter point for hnsw3 *L* ← level of *ep* // top layer for *hnsw*

 $4 l \leftarrow |-\ln(unif(0..1)) \cdot m_L| // \text{ new element's level}$ 5 for $l_c \leftarrow L \dots l+1$

 $W \leftarrow SEARCH-LAYER(q, ep, ef=1, l_c)$ $ep \leftarrow \text{get the nearest element from } W \text{ to } q$

8 for $l_c \leftarrow \min(L, l) \dots 0$

 $W \leftarrow SEARCH-LAYER(q, ep, efConstruction, l_c)$ $neighbors \leftarrow SELECT-NEIGHBORS(q, W, M, l_c) // alg. 3 or alg. 4$

for each $e \in neighbors$ // shrink connections if needed $eConn \leftarrow neighbourhood(e)$ at layer l_c 13

if $|eConn| > M_{max}//$ shrink connections of e 14

// if $l_c = 0$ then $M_{max} = M_{max0}$

15 $eNewConn \leftarrow SELECT-NEIGHBORS(e, eConn, M_{max}, l_c)$

// alg. 3 or alg. 4 set neighbourhood(e) at layer lc to eNewConn

add bidirectionall connectionts from *neighbors* to *q* at layer *lc*

17 $ep \leftarrow W$ 18 if l > L14

set enter point for hnsw to q

Candidate Selection Simple

Q: Which *M* nodes to take out of *efConstruction* candidates?

A: Naive way – take M closest candidates

Then a node X is inserted into the graph and needs to be linked to M=2 other vertices: B and C.

However, ideally it can be better for navigation if the region A and B can be connected.

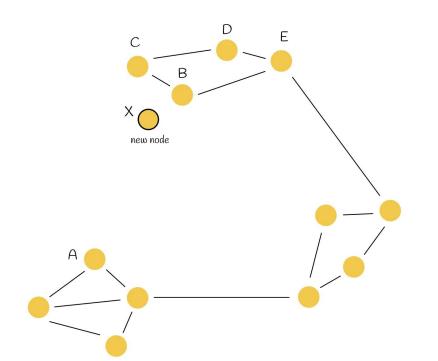
Algorithm 3

SELECT-NEIGHBORS-SIMPLE(q, C, M)

Input: base element q, candidate elements C, number of neighbors to return M

Output: *M* nearest elements to *q*

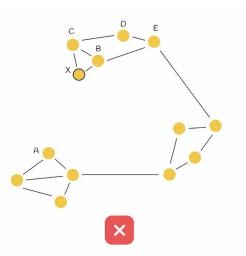
return *M* nearest elements from *C* to *q*

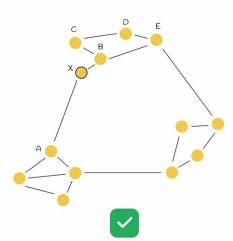


Candidate Selection Heuristic

The heuristic considers both:

- The closest distances between nodes
- The connectivity of different regions on the graph





Algorithm 4

SELECT-NEIGHBORS-HEURISTIC(q, C, M, lc, extendCandidates, keep-PrunedConnections)

Input: base element *q*, candidate elements *C*, number of neighbors to return *M*, layer number *lc*, flag indicating whether or not to extend candidate list *extendCandidates*, flag indicating whether or not to add discarded elements *keepPrunedConnections*

Output: *M* elements selected by the heuristic

- $1 R \leftarrow \emptyset$
- 2 $W \leftarrow C$ // working queue for the candidates
- 3 if extendCandidates // extend candidates by their neighbors
- 4 **for** each $e \in C$
- for each $e_{adj} \in neighbourhood(e)$ at layer l_c
- 6 **if** eadj ∉ W
- 7 $W \leftarrow W \cup e_{adj}$
- 8 $W_d \leftarrow \emptyset$ // queue for the discarded candidates
- 9 while |W| > 0 and |R| < M
- 10 $e \leftarrow$ extract nearest element from W to q
- 11 **if** e is closer to q compared to any element from R
- 12 $R \leftarrow R \cup e$
- 13 else
- 14 $W_d \leftarrow W_d \cup e$
- 15 **if** keepPrunedConnections // add some of the discarded // connections from W_d
- 16 while $|W_d| > 0$ and |R| < M
- 17 $R \leftarrow R \cup \text{extract nearest element from } W_d \text{ to } q$
- 18 return R

Complexity Analysis

Search takes *O(logn)* time in total

Insertion of a single vertex: O(logn)

HNSW construction requires O(n * logn) time in total

Evaluation - Implementation

 HNSW implementation uses custom distance functions together with C-style memory management.

Utilized nmslib implementation of sw-graph for NSW.

Compare with the most up-to-date SOTA.

Compare with the SOTA in Euclid Spaces with open-source implementation.

Evaluation - Method

Comparison with Baseline NSW

Comparison in Euclid Spaces

Comparison in General Space

Comparison with product quantization based algorithms.

Evaluation - HNSW vs. Baseline NSW

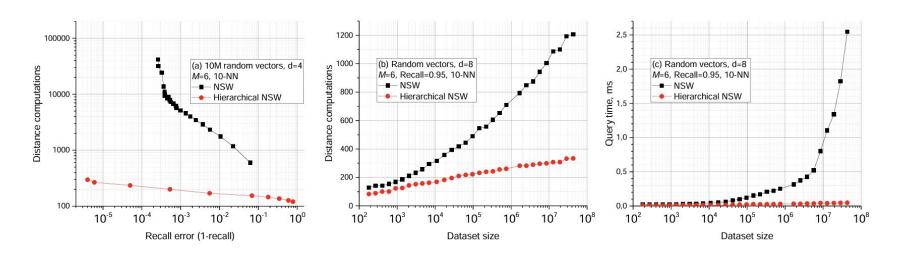


Fig. 12. Comparison between NSW and Hierarchical NSW: (a) distance calculation number vs accuracy tradeoff for a 10 million 4-dimensional random vectors dataset; (b-c) performance scaling in terms of number of distance calculations (b) and raw query(c) time on a 8-dimensional random vectors dataset.

Evaluation - Euclid Spaces - Algorithms to Compare

- Baseline NSW Algorithm
- FLANN
- Annoy
- VP-tree
- FALCONN

Evaluation - Euclid Spaces - Datasets

TABLE 1
Parameters of the used datasets on vector spaces benchmark.

Dataset	Description	Size	d	BF time	Space
SIFT	Image feature vectors [13]	1M	128	94 ms	L ₂
GloVe	Word embeddings trained on tweets [52] 1.2M 100 95 ms				cosine
CoPhIR	MPEG-7 features extracted from the images [53]	2M	272	370 ms	L ₂
Random vectors	Random vectors in hypercube	30M	4	590 ms	L ₂
DEEP	One million subset of the billion deep image features dataset [14]		96	60 ms	L ₂
MNIST	NIST Handwritten digit images [54]			22 ms	L ₂

Evaluation - Euclid Spaces

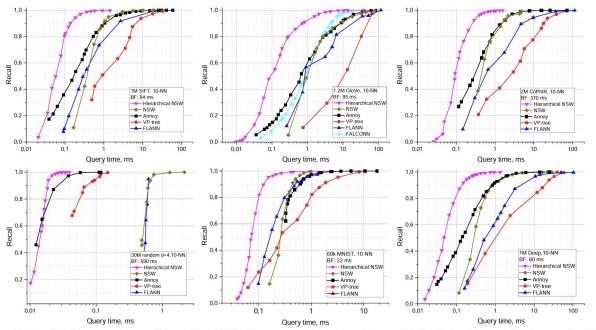


Fig. 13. Results of the comparison of Hierarchical NSW with open source implementations of K-ANNS algorithms on five datasets for 10-NN searches. The time of a brute-force search is denoted as the BF.

Evaluation - General Spaces - Purpose & Algorithms

- Baseline NSW algorithm has several problems on low dimensional datasets as suggested in the paper "Permutation search methods are efficient, yet faster search is possible."
- VP-tree
- Permutation Techniques (NAPP & Brute Force Filtering)
- Baseline NSW Algorithm
- NNDescent-produced proximity graphs

Evaluation - General Spaces - Datasets

TABLE 2.
Used datasets for repetition of the Non-Metric data tests subset.

Dataset	Description		d	BF time	Distance	
Wiki-sparse	TF-IDF (term frequency–inverse document frequency) vectors (created via GENSIM [58])	4M	105	5.9 s	Sparse cosine	
Wiki-8	Topic histograms created from sparse TF-IDF vectors of the wiki-sparse dataset (created via GENSIM [58])	2M	8		Jensen- Shannon (JS) divergence	
Wiki-128	Topic histograms created from sparse TF-IDF vectors of the wiki-sparse dataset (created via GENSIM [58])	2M	128	1.17 s	Jensen- Shannon (JS) divergence	
ImageNet	Signatures extracted from LSVRC-2014 with SQFD (signature quadratic form) distance [59]	1M	272	18.3 s	SQFD	
DNA	DNA (deoxyribonucleic acid) dataset sampled from the Human Genome 5 [34].	1M	-	2.4 s	Levenshtein	

Evaluation - General Spaces

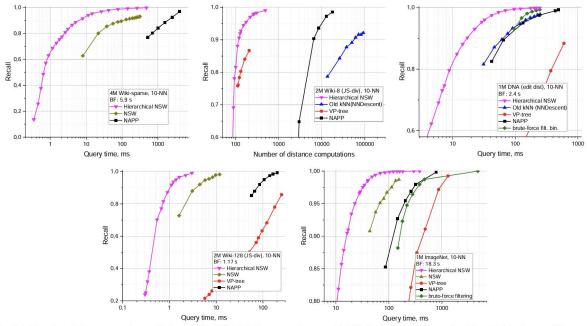


Fig. 14. Results of the comparison of Hierarchical NSW with general space K-ANNS algorithms from the Non Metric Space Library on five datasets for 10-NN searches. The time of a brute-force search is denoted as the BF.

Evaluation - HNSW vs product quantization based algorithms

- PQ-Algorithm: SOTA on billion scale datasets.
- Compare HNSW with SOTA PQ Algorithm in the library: Faiss.

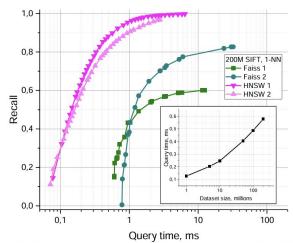


Fig. 15 Results of comparison with Faiss library on the 200M SIFT dataset from [13]. The inset shows the scaling of the query time vs the dataset size for Hierarchical NSW.

TABLE 3.

Parameters for comparison between Hierarchical NSW and Faiss on a 200M subset of 1B SIFT dataset.

Algorithm	Build time	Peak memory (runtime)	Parameters
Hierarchical NSW	5.6 hours	64 Gb	M=16, efConstruction=500 (1)
Hierarchical NSW	42 minutes	64 Gb	M=16, efConstruction=40 (2)
Faiss	12 hours	30 Gb	OPQ64, IMI2x14, PQ64 (1)
Faiss	11 hours	23.5 Gb	OPQ32, IMI2x14, PQ32 (2)

Conclusion

 HNSW provides a groundbreaking approach to nearest neighbor search, balancing speed and accuracy effectively even in challenging, high-dimensional spaces.

 The HNSW graph demonstrates robustness to various dataset that was not solvable by baseline NSW. It maintains good performance across different types of datasets without significant tradeoffs.

 This method sets a new benchmark for nearest neighbor searches, offering significant implications for machine learning and data retrieval.

Limitations

 Constructing and maintaining the HNSW graph can consume significant memory, especially for large datasets. This can limit the scalability of the method on memory-constrained systems or for applications with extremely large datasets.

 The search in the HNSW structure always starts from the top layer, thus the structure cannot be easily made distributed like baseline NSW.

Future Work

 The number of added connections per layer M can be a meaningful parameter to tune that strongly affects the construction of the index, thus might improve efficiency and effectiveness of HNSW.

 It would also be interesting to compare HNSW on the full 1B SIFT and 1B DEEP datasets and with functionalities such as element updates and removal.

Design a distributed pipeline for speedup and memory optimization.

Thanks!