

IR From Bag-of-words to BERT and Beyond through Practical Experiments

A Search Solutions 2022 tutorial with PyTerrier

Starting at 14:45 GMT

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(*Alphabetical ordering)

PART 3

INDEX AUGMENTATION &

LEARNED SPARSE RETRIEVAL &

DENSE RETRIEVAL



Intended Learning Outcomes



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Part 3 – Understand the most recent effective retrieval architectures that learn representations of documents, both in leveraging traditional index structures (index augmentation & sparse retrieval), as well as new similarity search systems (dense retrieval)

ILO 3A. Understand neural augmentation models, such as Doc2Query

ILO 3B Understand learned sparse retrieval methods, such as DeepCT, DeeplImpact and SPLADE

ILO 3C. Understand dense retrieval models and how embedding indexes work, such as FAISS, ANCE and ColBERT

ILO 3D. Perform experiments with ANCE and ColBERT in a Python notebook.

Outline of Part 3



Part 3A: Neural Index Augmentation

Part 3B: Learned Sparse Retrieval

Part 3C: Dense Retrieval

Part 3D: Embedding Indexes and Vector Search

Part 3A

NEURAL INDEX AUGMENTATION

Document Augmentation



E.g., Inverted Index

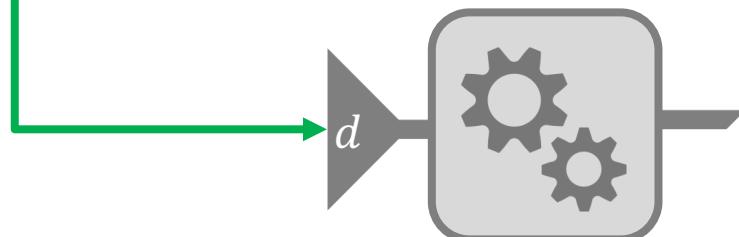
docno	text
ug7v899j	OBJECTIVE: This retrospective chart review describes the...
02tnwd4m	Inflammatory diseases of the respiratory tract are common...
ejv2xln0	Endothelin-1 (ET-1) is a 21 amino acid peptide with diverse...
...	...

Document Augmentation



E.g., Inverted Index

docno	text
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...	...

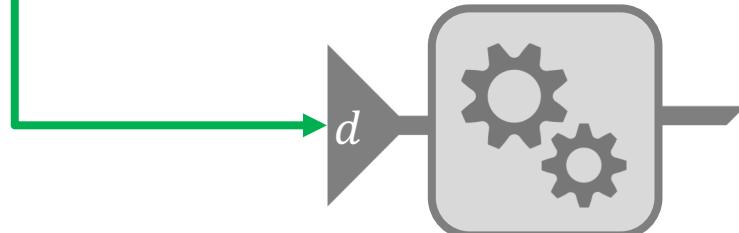


where is mycoplasma p where is
jeddah hospital located what
culture proves mycoplasma
pneumoniae

Document Augmentation



docno	text	newfield
ug7v899j	OBJECTIVE: This retrospective chart review describes the...	where is mycoplasma p where is jeddah...
02tnwd4m	Inflammatory diseases of the respiratory tract are common...	
ejv2xln0	Endothelin-1 (ET-1) is a 21 amino acid peptide with diverse...	
...	...	

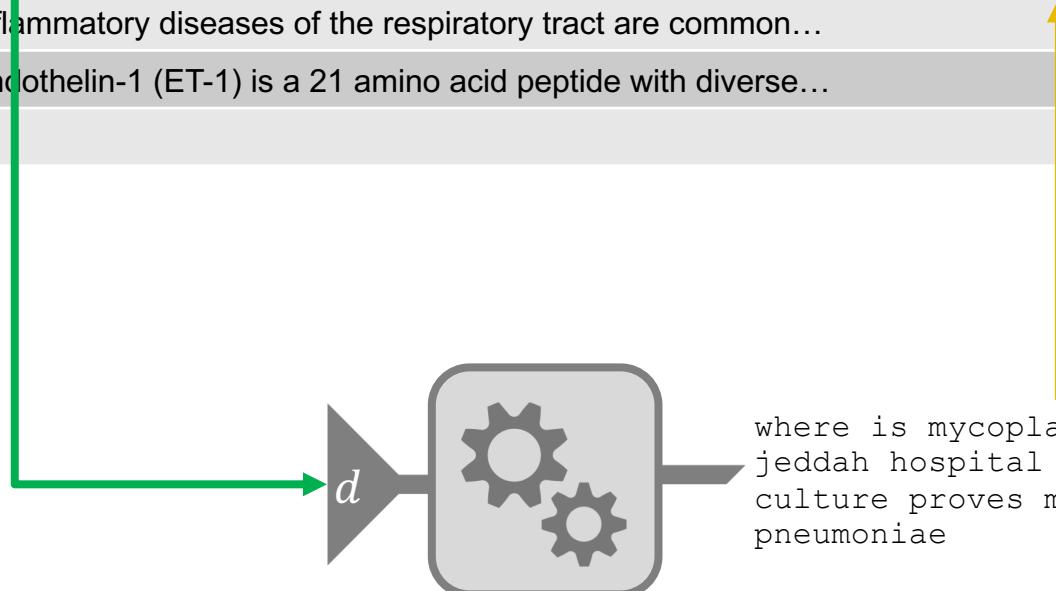


where is mycoplasma p where is jeddah hospital located what culture proves mycoplasma pneumoniae

Document Augmentation



docno	text
ug7v899j	OBJECTIVE: This retrospective chart review describes the... where is mycoplasma p where is jeddah...
02tnwd4m	Inflammatory diseases of the respiratory tract are common...
ejv2xln0	Endothelin-1 (ET-1) is a 21 amino acid peptide with diverse...
...	...



Main idea: Use a causal language model to generate additional text to add to documents when indexing.
At retrieval time, use standard retrieval model (e.g., BM25).

How are the queries generated?

Abstract:

Nidovirus subgenomic mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous subgenomic RNA synthesis that resembles copy-choice RNA recombination. During this process, the nascent RNA strand is transferred from one site in the template to another, during either plus or minus...

Generated Queries:

- where does mrna originate
- where does subgenomic rna come from
- where is the nidovirus mrna?
- when a nidovirus is produced, its genome is quizlet
- what is nidovirus mrna
- where in a nidovirus can one mrna be derived
- where is the leader sequence derived
- what types of mrna are found in nidovirus
- where are the nidovirus genomes derived from
- ...



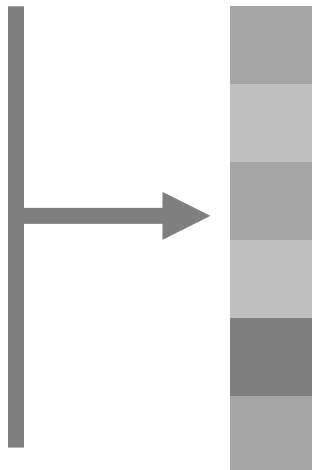
Training: optimize to generate queries from passages on a collection with many queries (e.g., MS-MARCO)

Most recent version: Use T5 model for generation (docTTTTquery)

Doc2Query

Abstract:

Nidovirus subgenomic mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous subgenomic RNA synthesis that resembles copy-choice RNA recombination. During this process, the nascent RNA strand is transferred from one site in the template to another, during either plus or minus...

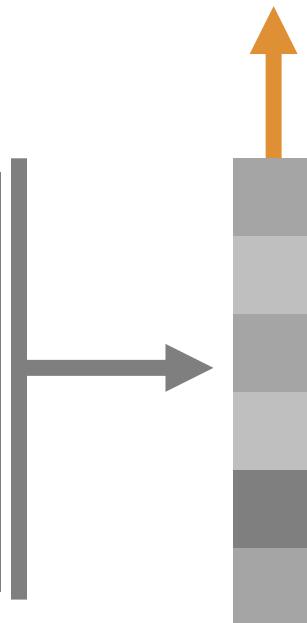


Step 1: Source text encoded
e.g., via RNN or Transformer

Doc2Query

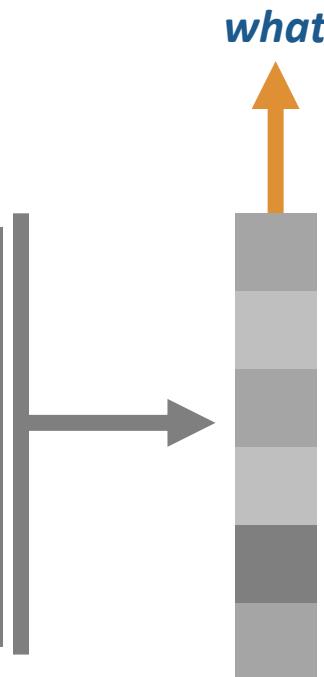
Abstract:

Nidovirus subgenomic mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous subgenomic RNA synthesis that resembles copy-choice RNA recombination. During this process, the nascent RNA strand is transferred from one site in the template to another, during either plus or minus...



Step 2: Text iteratively generated from encoded document
e.g., via RNN or transformer, using beam search

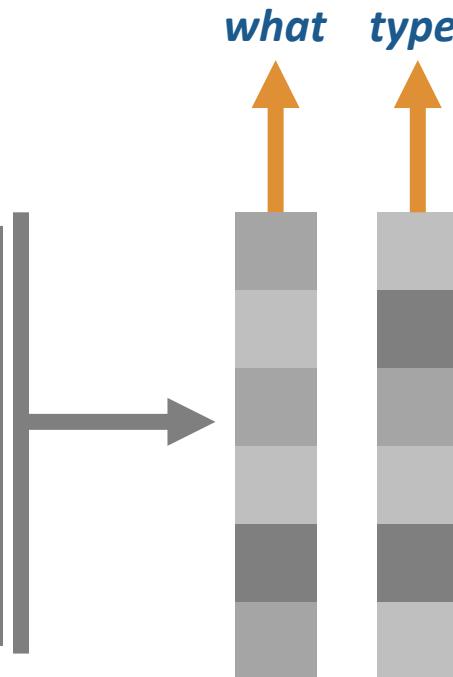
Abstract:
Nidovirus subgenomic mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous subgenomic RNA synthesis that resembles copy-choice RNA recombination. During this process, the nascent RNA strand is transferred from one site in the template to another, during either plus or minus...



Step 2: Text iteratively generated from encoded document
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Abstract:

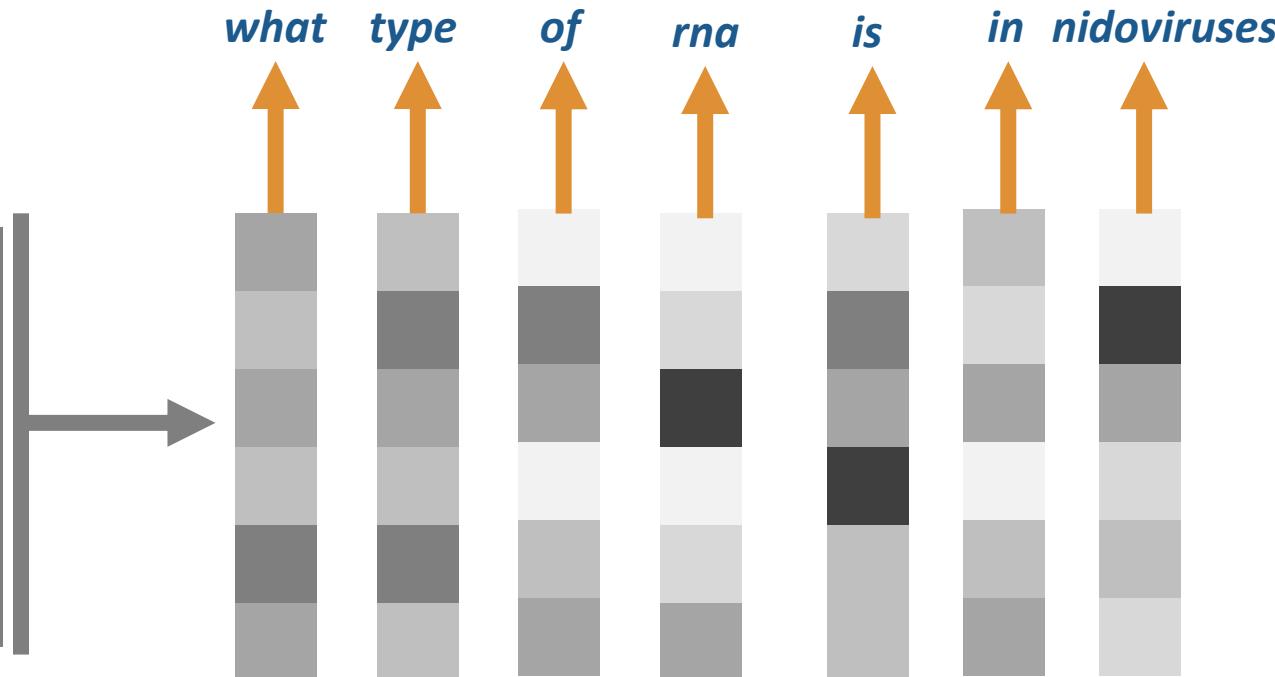
Nidovirus subgenomic mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous subgenomic RNA synthesis that resembles copy-choice RNA recombination. During this process, the nascent RNA strand is transferred from one site in the template to another, during either plus or minus...



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Step 2: Text iteratively generated from encoded document
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Doc2Query Practical

```
from pyterrier_doc2query import Doc2Query
doc2query = Doc2Query(out_attr="text", batch_size=8)
```

Example doc2query outputs

```
df.iloc[0]['text']
```

'OBJECTIVE: This retrospective chart review describes the epidemiology and clinical features of 40 patients with culture-proven Mycoplasma pneumoniae infections at King Abdulaziz University Hospital, Jeddah, Saudi Arabia. MET HODS: Patients with positive M. pneumoniae cultures from respiratory specim

```
doc2query.transform(df).iloc[0]['text']
```

'what is culture confirmed mycoplasma where is mycoplasma pneumonia located in saudi arabia? where is mycoplasma cultured'



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Part 3B

LEARNED SPARSE RETRIEVAL

Learned Sparse Retrieval

Main question: Can we leverage the neural IR effectiveness improvements together with the sparse retrieval efficiency in query processing?

- In sparse retrieval, documents are stored in an inverted index data structure
- The inverted index is highly specialized toward efficient query processing
- Major data structure in search engines
- In neural IR, semantic models outperform lexical models in ad-hoc search tasks
- Storing semantic information is expensive
- Processing semantic information is expensive

Main idea: Boost the term frequency of important words by repeating them in the text.

Abstract:

Nidovirus subgenomic mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous subgenomic RNA synthesis that resembles copy-choice RNA recombination. During this process, the nascent RNA strand is transferred from one site in the template to another, during either plus or minus...



Abstract:

Nidovirus Nidovirus Nidovirus subgenomic subgenomic subgenomic mRNAs mRNAs mRNAs contain a leader sequence derived from the 5' end of the genome fused to different sequences ('bodies') derived from the 3' end. Their generation involves a unique mechanism of discontinuous **subgenomic subgenomic RNA RNA** synthesis that resembles copy-choice **RNA RNA RNA** recombination. During this process, the nascent **RNA RNA RNA** strand is transferred from one site in the template to another, during either plus or minus...



Training: Predict the terms that match relevant query terms.

DeepCT practical

```
deepct = pyterrier_deepct.DeepCTTransformer(  
    "bert-base-uncased/bert_config.json",  
    "marco/model.ckpt-65816"  
)  
deepct(pd.DataFrame([  
    {'docno': '0', 'text': '''The presence of communication amid scientific minds  
    was equally important to the success of the Manhattan  
    Project as scientific intellect was. The only cloud  
    hanging over the impressive achievement of the atomic  
    researchers and engineers is what their success truly  
    meant; hundreds of thousands of innocent lives  
    obliterated.'''  
}]))
```

presence communication scientific minds equally important to success manhattan
project intellect cloud impressive achievement atomic researchers engineers truly
meant lives obliterated

Main idea: Compute a quantized score (**impact**) to store in posting list

- Not just BM25 + learned TF as in DeepCT
- Add text to documents using a contextualized language model
- Address the vocabulary mismatch problem
- Exploit BERT to compute a single score for each query-document pair
- Quantize linearly the scores into positive integers
- Store quantized scores as frequencies in postings, using simple sum as weighting model

DeepImpact practical

- Sparse Indexing – Learned

```
vaswani = pt.get_dataset("vaswani")

pt_index_path = './terrier_di_vaswani'

parent = pt.index.IterDictIndexer(pt_index_path)
parent.setProperty("termpipelines", "")

indexer = DeepImpactIndexer(parent, batch_size=32)
indexer.index(vaswani.get_corpus_iter())

index_ref = pt.IndexRef.of(pt_index_path + "/data.properties")
index_di = pt.IndexFactory.of(index_ref)
```

```
pt_index_path = './terrier_vaswani'

indexer = pt.index.IterDictIndexer(pt_index_path)
indexer.setProperty("termpipelines", "")
index_ref = indexer.index(vaswani.get_corpus_iter())

index_ref = pt.IndexRef.of(pt_index_path + "/data.properties")
index = pt.IndexFactory.of(index_ref)
```

- End-to-end Retrieval

```
pt.Experiment([
    pt.BatchRetrieve(index, wmodel="BM25"),
    pt.BatchRetrieve(index_di, wmodel="Tf")
],
vaswani.get_topics(), vaswani.get_qrels(),
names=['bm25', "deep_impact"],
eval_metrics=["map", "recip_rank", "ndcg_cut_10"]
)
```

	name	map	recip_rank	ndcg_cut_10
0	bm25	0.143471	0.516597	0.249405
1	deep_impact	0.186312	0.661398	0.332272

Original Terms: of the in origin atmosphere upper nitrogen ionization
DeepImpact Terms: of the a in and to is described are design suitable discussed
characteristics probe positive small such experiments density measurement
ionosphere type designed rocket spherical laboratory ion borne investigate
adopted meshed

Main idea: compute scores and query/document expansions at the same time (cf. EPIC)

- BERT models compute a term language model for each output embeddings
 - A language model is a probability distribution over the vocabulary
 - These language models capture the semantic correlations between the input term and all other terms in the vocabulary
 - They can be used to expand documents with new terms or compress them by removing existing terms

Main idea: compute scores and query/document expansions at the same time (cfr. EPIC)

- Compute a (un-normalized) LM for each token in the document
- Filter the weights with logarithm to saturate high values
- Sum everything up to get a document-level language model
- Use the highest-valued components to add new terms to the documents
 - Special training to boost sparsity (FLOPS regularisation)

SPLADE practical

- Sparse Indexing – Learned

```
1 import pandas as pd
2 import pyterrier as pt ; pt.init()
3 import pyt_splade

1 splade = pyt_splade.SpladeFactory(agg='max')

1 doc_pipeline = splade.indexing()

1 doc_pipeline(pd.DataFrame([
2     {'docno': '0', 'text': 'The presence of communica
3     {'docno': '100', 'text': "Antonín Dvorák (1841-19
4     {'docno': '1000', 'text': 'QuickFacts Matanuska-S
5 ]))
```



the presence of communication amid scientific minds was equally important to the success of the manhattan project as scientific intellect was . the only cloud hanging over the impressive achievement of the atomic researchers and engineers is what their success truly meant ; hundreds of thousands of innocent lives ob ##lite ##rated .

Expansion Tokens: communications importance successful intellectual intelligence mind clouds projects achievements york amongst engineer bomb among engineering atom scientist helped nuclear amidst thinking impact thought destroyed scientists experiment existence telecommunications many failed because research thoughts hang message researcher science millions life death die contributed brains mean enlightenment murder minded significance ##rate survival hiroshima radio significant leader tunnel threat were clouded goal role really invented ##athic accomplishments controversial successes ended resulted purpose defeated casualties interesting physicist amazing communicate roosevelt people intent discovery started franklin telegraph within einstein journalist merit lack army tech hoped mental truman ny ##bility rocket led story great ##bm went study suspended respect comet radios collaboration alla destruction kennedy perception died committee studies society also said radioactive relativity conducted spirit results miracle suicide necessary crash result thinkers affect war ##rut absence contribution mission forensic news destroy knowledge wanted involved

SPLADE practical

- Indexing & End-to-end Retrieval on MSMarco v1

```
import pyterrier as pt
pt.init(version='snapshot')
import pyt_splade

dataset = pt.get_dataset('irds:msmarco-passage')
splade = pyt_splade.SpladeFactory()

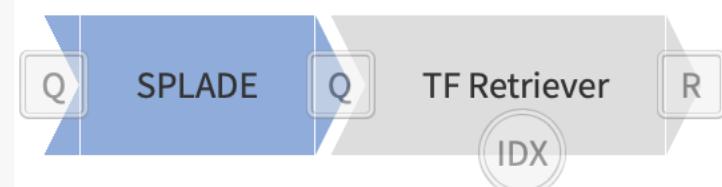
indexer = pt.IterDictIndexer('./msmarco_psg', pretokenized=True)

idxer_pipe = splade.indexing() >> indexer
idxer_pipe.index(dataset.get_corpus_iter())
```



```
bm25_retr = pt.BatchRetrieve.from_dataset('msmarco_passage', 'terrier_stemmed', wmodel='BM25')
splade_retr = splade.query() >> pt.BatchRetrieve('./msmarco_psg', wmodel='Tf')

pt.Experiment(
    [splade_retr, bm25_retr],
    pt.get_dataset('msmarco_passage').get_topics('test-2019'),
    pt.get_dataset('msmarco_passage').get_qrels('test-2019'),
    batch_size=200,
    filter_by_qrels=True,
    eval_metrics=[RR(rel=2), nDCG@10, nDCG@100, AP(rel=2)],
    names=['splade', 'bm25_stemmed'])
```



name	RR(rel=2)	nDCG@10	nDCG@100	AP(rel=2)
splade	0.918605	0.730178	0.671292	0.503885
bm25_stemmed	0.641565	0.47954	0.487416	0.286448

Other Approaches

Method	Query	Passage	Reg.	Supervision		
				Level	Neg.	Type
DeepCT [3]	MLP	MLP	-	Term	-	-
uniCOIL [14]	MLP	MLP	-	Passage	BM25(s)	Human
uniCOIL _{dT5q} [14]	MLP	expMLP	-	Passage	BM25(s)	Human
uniCOIL _{tilde} [14]	MLP	expMLP	-	Passage	BM25(s)	Human
EPIC [17]	MLP	MLM	Top-k	Passage	BM25	Human
DeepImpact [18]	MLP	expMLP	-	Passage	BM25	Human
TILDE [30]	BINARY	clsMLM	-	Term	-	-
TILDEv2 [30]	BINARY	expMLP	-	Passage	BM25(s)	Human
Sparta [29]	BINARY	MLM	-	Passage	BM25	Human
SPLADE-max [4]	MLM	MLM	FLOPs	Passage	BM25(s)	Human
DistilSPLADE-max [4]	MLM	MLM	FLOPs	Passage	Hard	Teacher

Summary

- **DeepCT**: use BERT to learn in-document term frequencies, to use with the classical BM25 model
- **DeeplImpact**: use BERT to learn in-document term scores, to store directly in the inverted index in quantised format
- **SPLADE**: use BERT to learn in-document term scores, to store directly in the inverted index, and to perform document expansion at indexing time and query expansion at query processing time
- Other approaches are variants of these approaches



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Part 3C

DENSE RETRIEVAL

Dense Retrieval

- Dense Retrieval, particularly for passages is of great interest
 - At indexing time, represent passages by one or more embeddings
 - At retrieval time, encode the query using the same model, and identify passages with embedding(s) **similar** to the query
- Great improvements have been proposed in large-scale **similarity search systems**
 - E.g. **FAISS** is the Facebook large scale nearest neighbour (NN) search system
 - FAISS supports for **exact** and **approximate search**
 - The type of search depends on the **problem dimension**
- In the following, we review the main dense retrieval models, then dense retrieval NN search

Sparse vs Dense Retrieval (I)

- **Sparse Retrieval**

- Every document is represented by a single score for each term in the lexicon V (equal to 0 for terms not appearing in the document)
- A document is a $|V|$ -dimensional vector with many 0s, i.e., sparse
- A query-document relevance score is computed as the sum of the score of the query terms in the documents
- Only query terms appearing in the document contributes to the final score
- A query is then a $|V|$ -dimensional vector with almost all entries set to 0, only a few set to 1

- **Pros**

- Efficient storage in inverted indexes
- Easily scalable
- 40+ years of query processing optimizations based on the inverted index
- Widely adopted as first stage ranker in cascading architectures

- **Cons**

- Very few relevance signals
- Not considering semantics information (except learned sparse models)
- Limited support for proximity signals

Sparse vs Dense Retrieval (II)

- **Dense Retrieval**

- Every document and query are represented by a D -dimension vector(s) of scores with few 0s, i.e., dense (embeddings)
- The query-document relevance score is computed as the L2 or cosine similarity between the query-document vectors

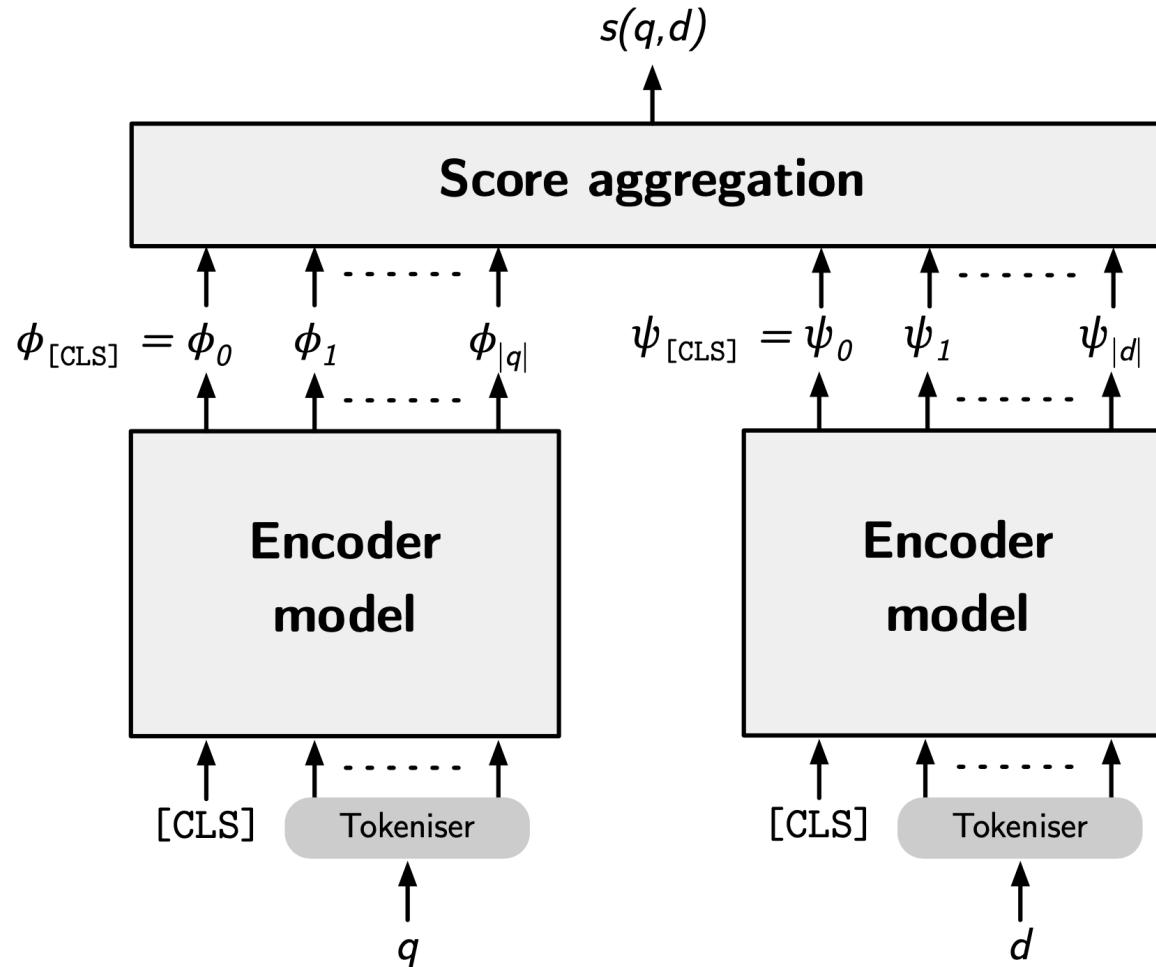
- **Pros**

- Efficient storage in similarity indexes
- 20+ years of query processing optimizations based on the nearest neighbour search
- Rich set of relevance signals
- Consider semantics

- **Cons**

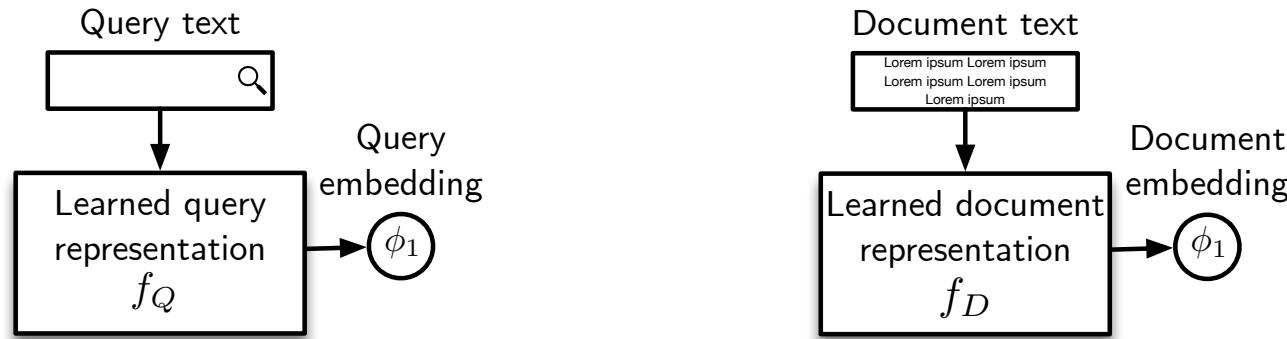
- Missing a clear interpretation of the embedding space
- Missing a clear interpretation of the captured semantics

Representation-focused System



Learning Representations

- Single Representation (USE, DPR, ANCE, STAR/ADORE)

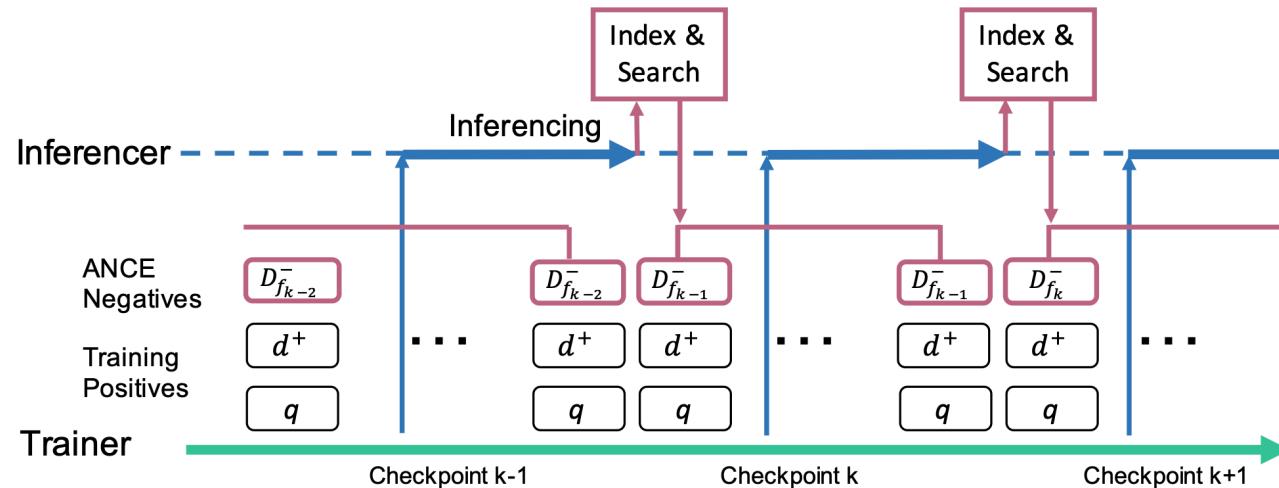
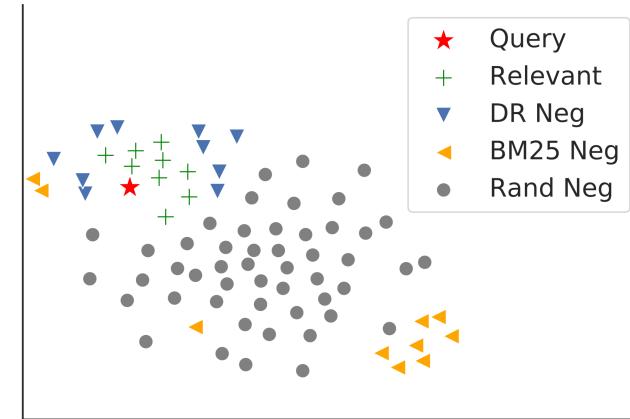


- Multiple Representation (Poly-encoders, ME-BERT, COIL, ColBERT)



ANCE: Approximate Negative Contrastive Learning

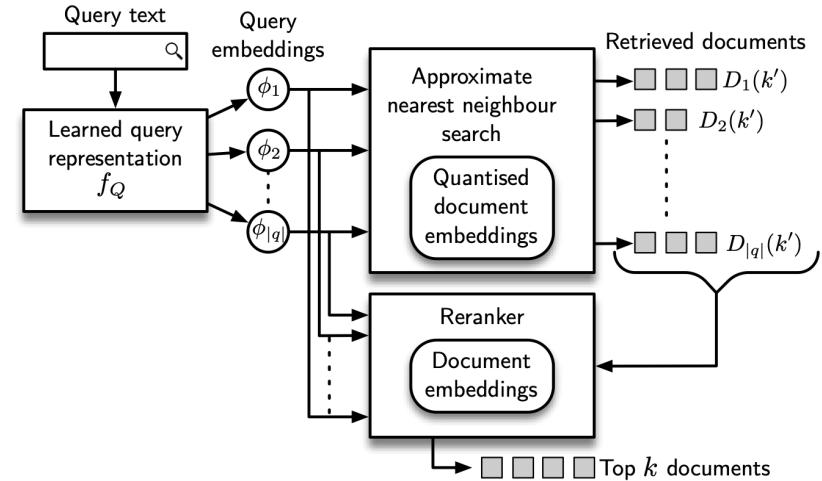
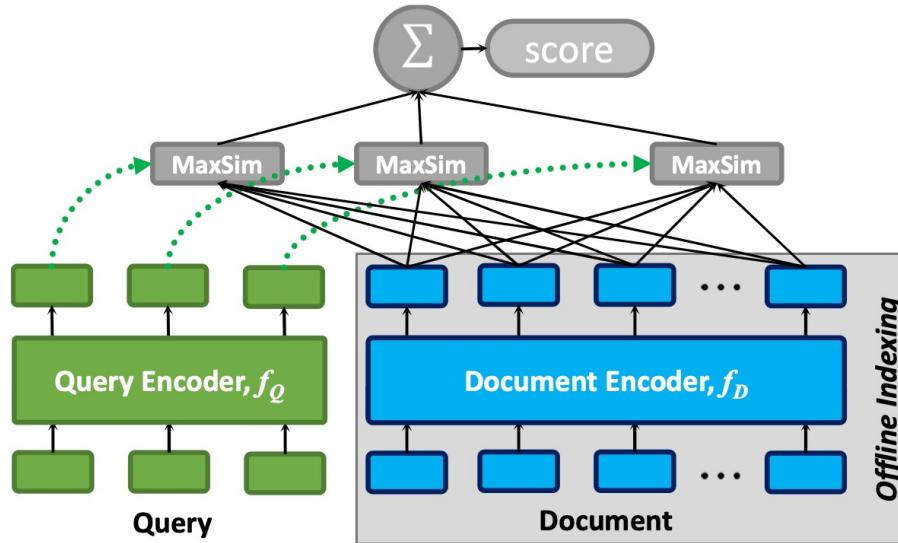
- Fully learnable representation
- Support for (approximate) NN search
- Problem: negative examples are usually drawn from a (classical) first stage
- How do we generate global negatives?
- Use the model learned so far → dynamic hard negatives



Exact NN Dense Retrieval (DPR, ANCE)

	MARCO Dev Passage Retrieval		TREC DL Passage NDCG@10	
	MRR@10	Recall@1k	Rerank	Retrieval
Sparse & Cascade IR				
BM25	0.240	0.814	—	0.506
Best DeepCT	0.243	n.a.	—	n.a.
Best TREC Trad Retrieval	0.240	n.a.	—	0.554
BERT Reranker	—	—	0.742	—
Dense Retrieval				
Rand Neg	0.261	0.949	0.605	0.552
NCE Neg	0.256	0.943	0.602	0.539
BM25 Neg	0.299	0.928	0.664	0.591
DPR (BM25 + Rand Neg)	0.311	0.952	0.653	0.600
BM25 → Rand	0.280	0.948	0.609	0.576
BM25 → NCE Neg	0.279	0.942	0.608	0.571
BM25 → BM25 + Rand	0.306	0.939	0.648	0.591
ANCE (FirstP)	0.330	0.959	0.677	0.648

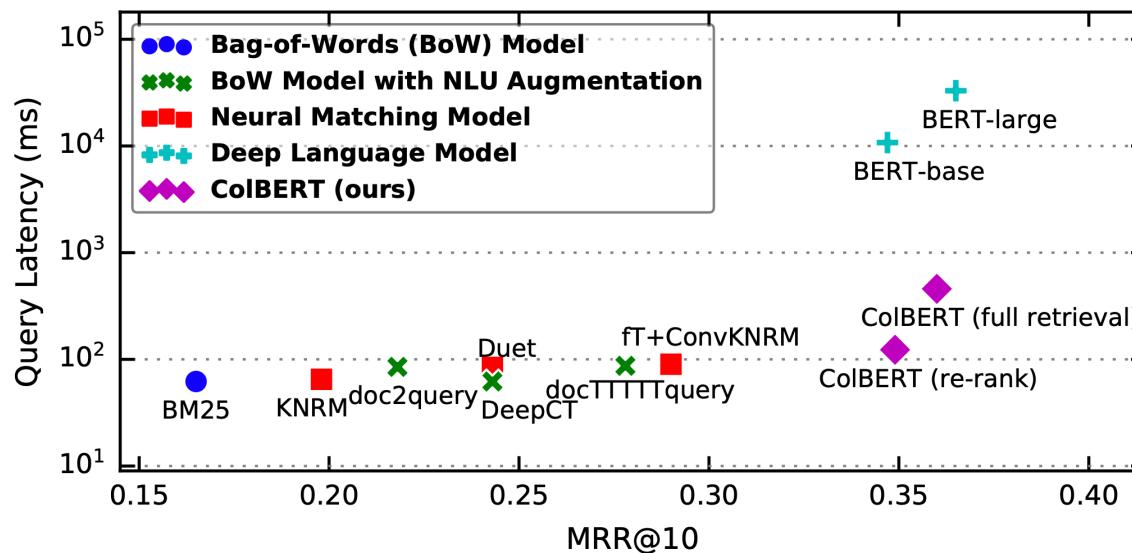
ColBERT: Contextualised Late Interaction BERT



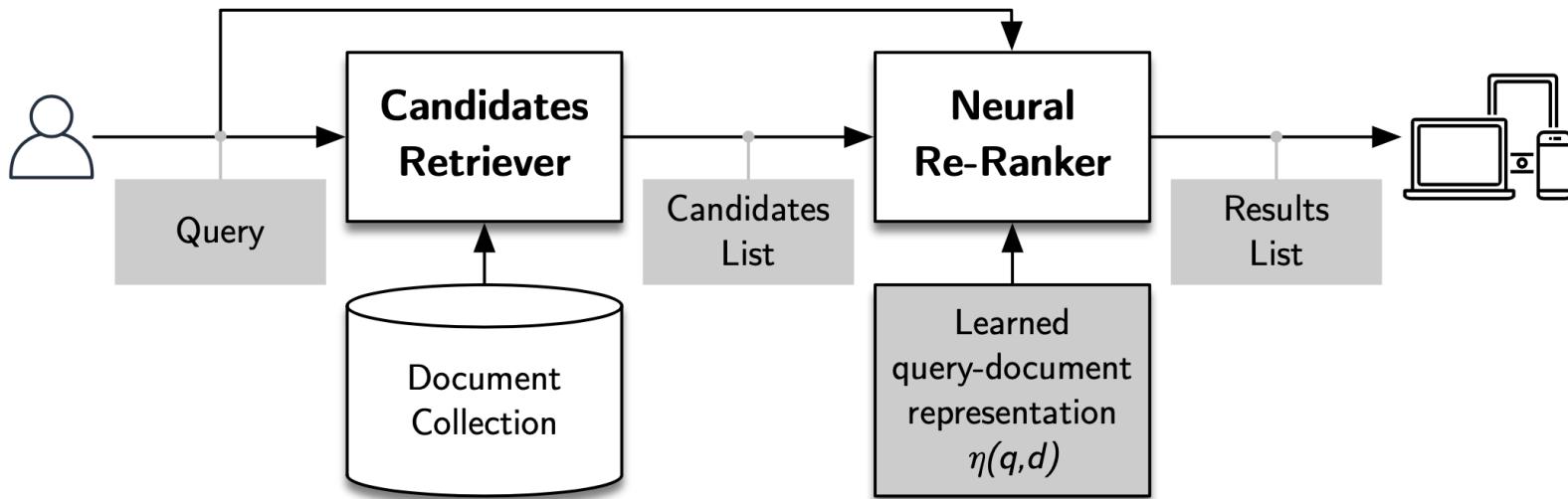
```
def end_to_end(self) -> TransformerBase:
    """
    Returns a transformer composition that uses a ColBERT FAISS index to retrieve documents, followed by a ColBERT index
    to perform accurate scoring of the retrieved documents. Equivalent to `colbertfactory.set_retrieve() >> colbertfactory.index_scorer()`.
    """
    #input: qid, query,
    #output: qid, query, docno, score
    return self.set_retrieve() >> self.index_scorer(query_encoded=True)
```

Approximate NN Dense Retrieval ColBERT

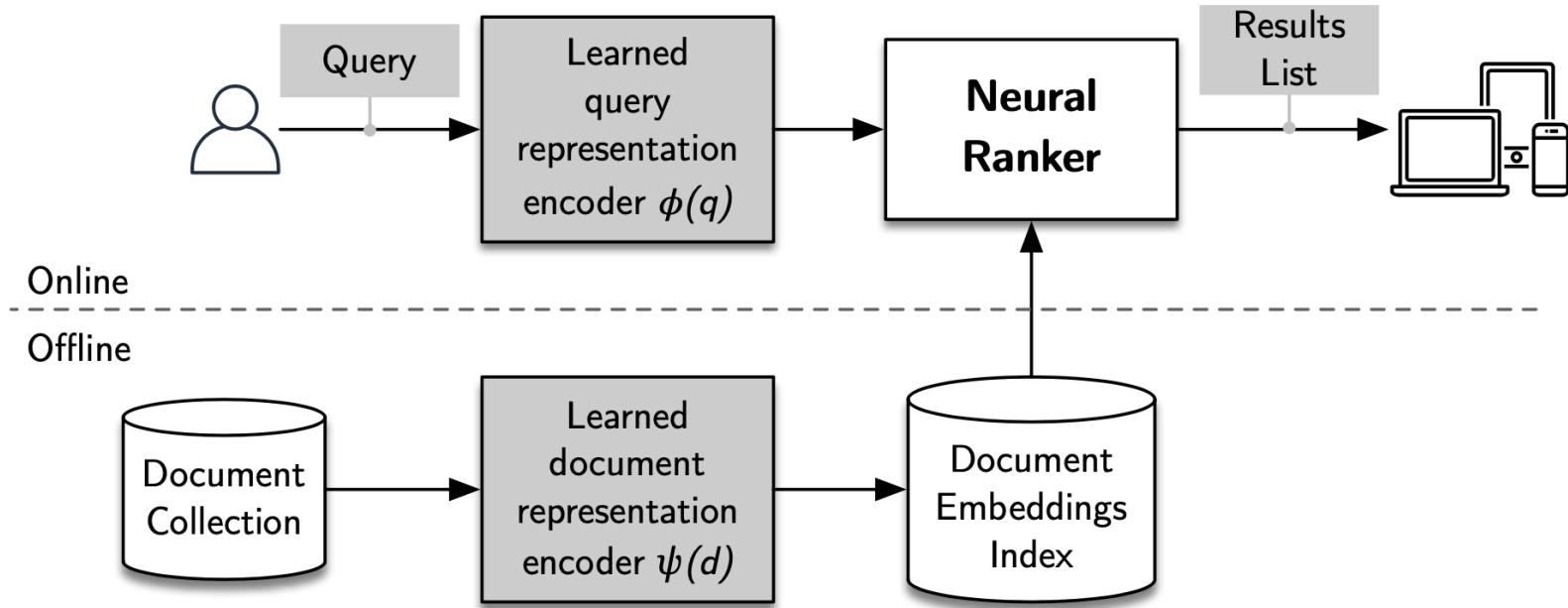
Method	MRR@10 (Dev)	MRR@10 (Local Eval)	Latency (ms)	Recall@50	Recall@200	Recall@1000
BM25 (official)	16.7	-	-	-	-	81.4
BM25 (Anserini)	18.7	19.5	62	59.2	73.8	85.7
doc2query	21.5	22.8	85	64.4	77.9	89.1
DeepCT	24.3	-	62 (est.)	69 [2]	82 [2]	91 [2]
docTTTTquery	27.7	28.4	87	75.6	86.9	94.7
ColBERT _{L2} (re-rank)	34.8	36.4	-	75.3	80.5	81.4
ColBERT _{L2} (end-to-end)	36.0	36.7	458	82.9	92.3	96.8



Re-ranking Pipeline for Interaction-focused Systems



Dense Retrieval for Representation-focused Systems

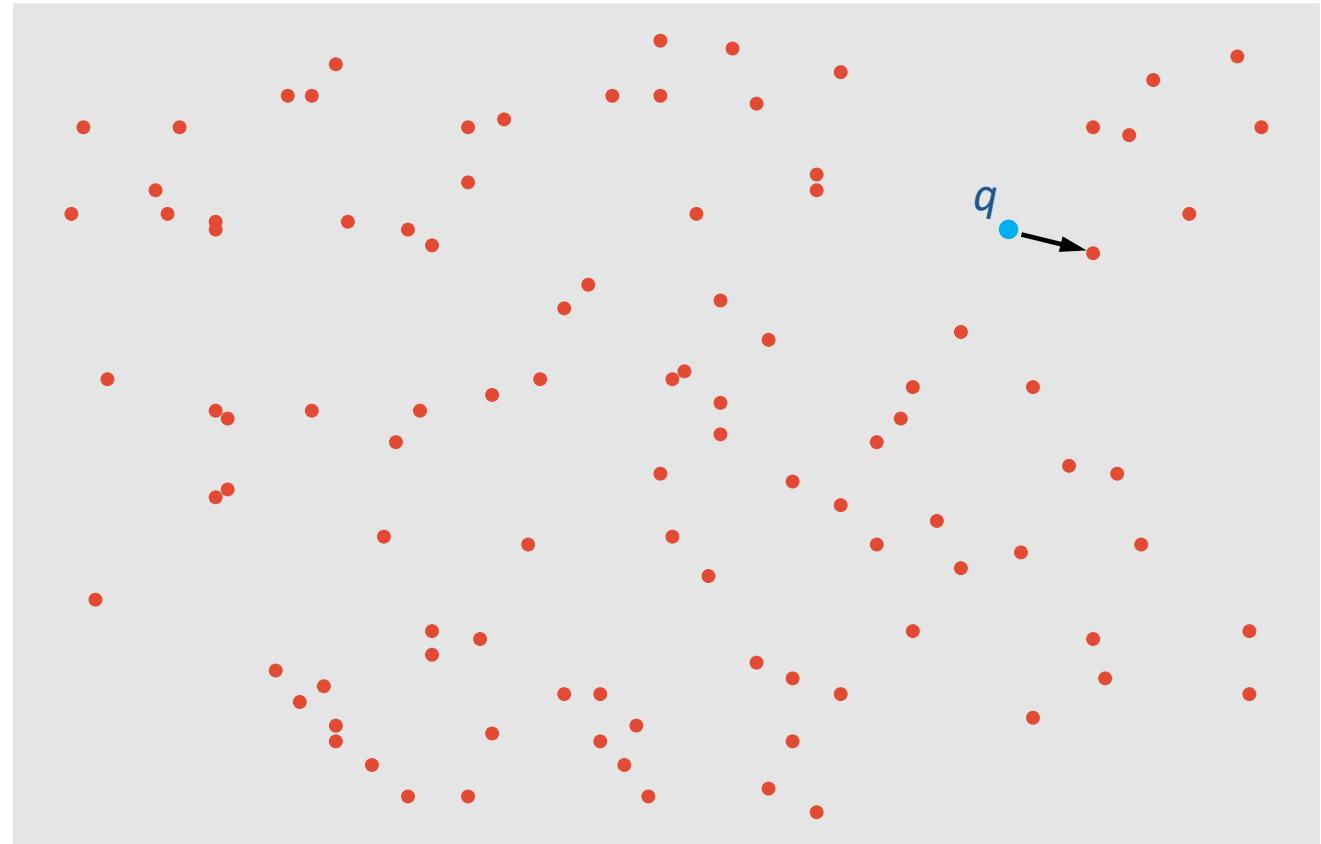


Part 3D

EMBEDDING INDEXES & VECTOR SEARCH

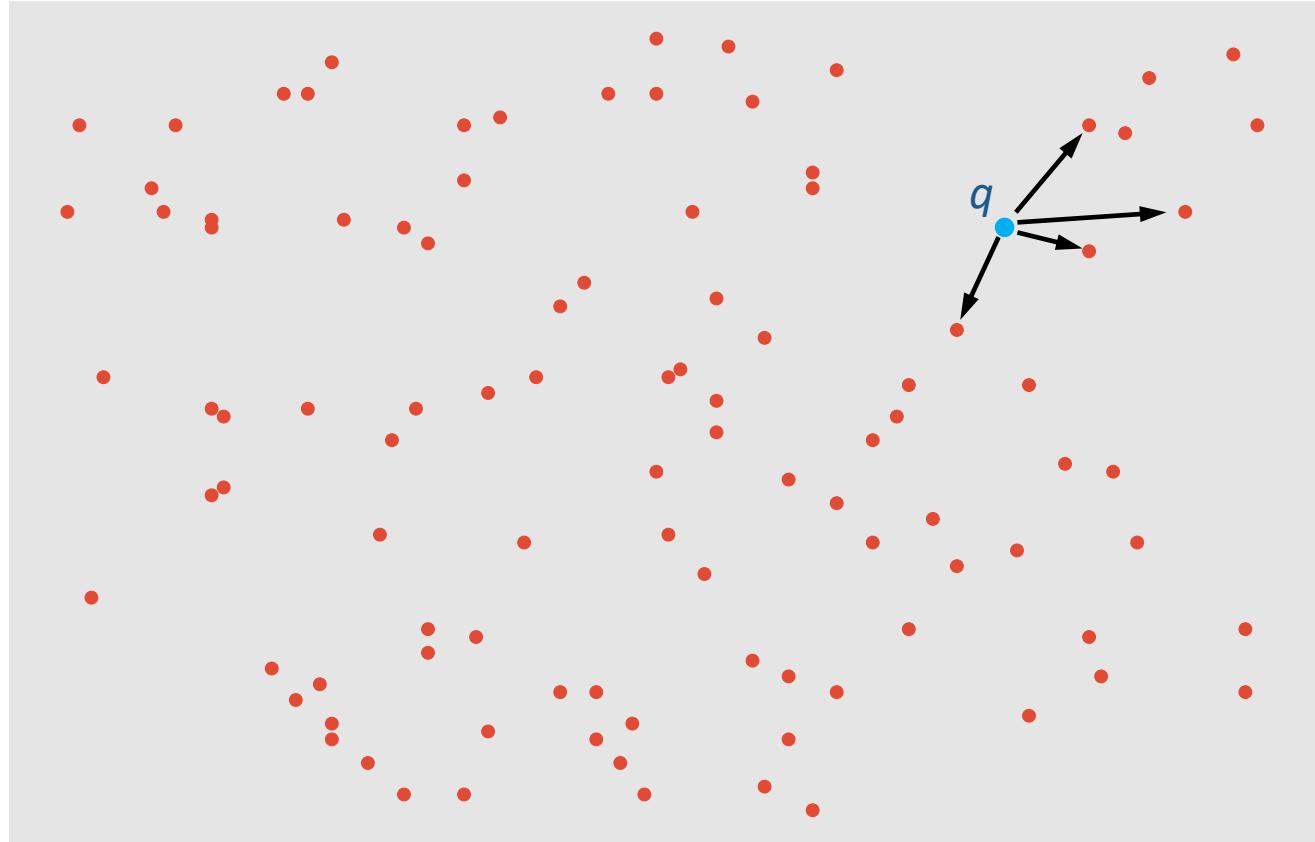
Nearest Neighbour

Complexity $O(n)$



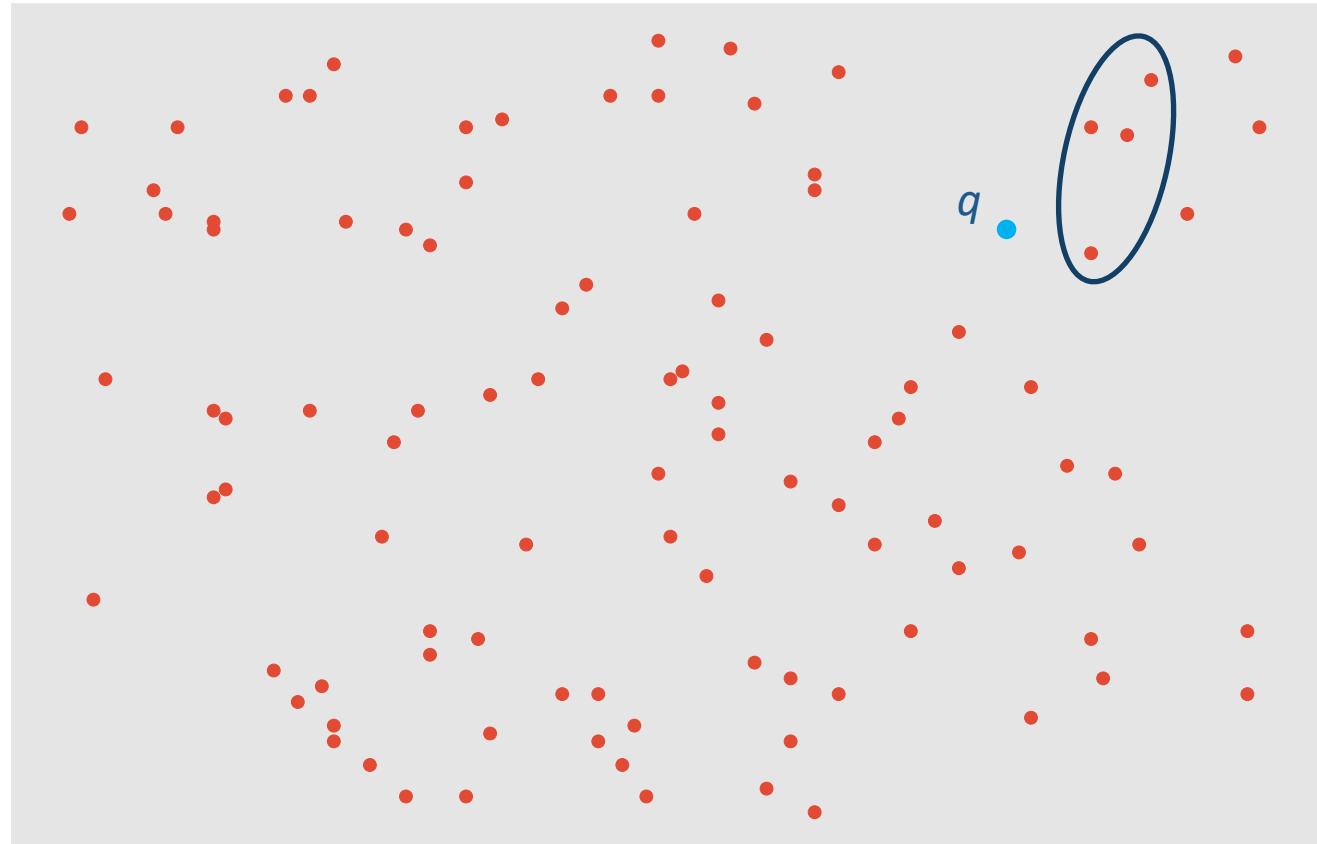
k Nearest Neighbours

Complexity $O(n \log k)$



Approximate Nearest Neighbours

- To make it **scalable**, we instead turn to **approximation** techniques
- **Avoid computing** the distance to every reference vector
- Return points **close** to the nearest neighbors
- Non-Exhaustive
- Common accuracy metric: recall@k



ANN Strategies

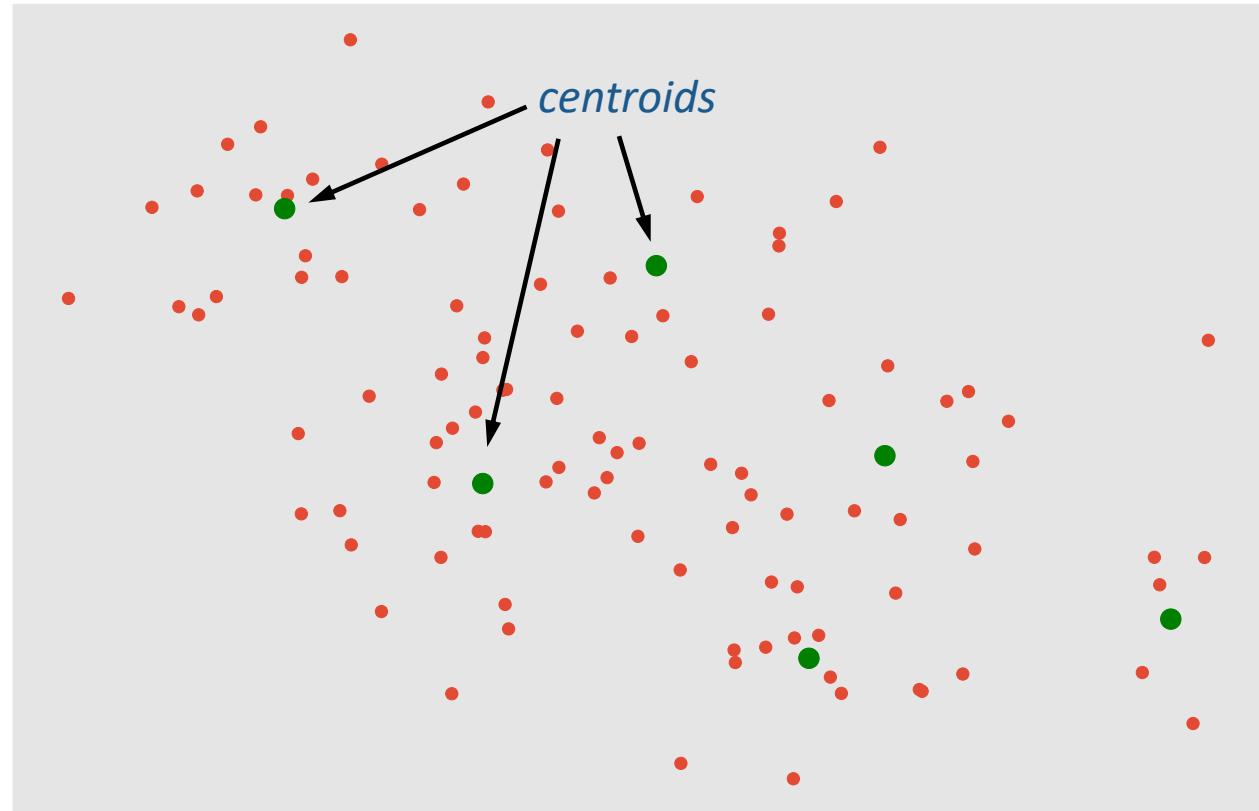


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- Trees and forests
- Locally Sensitive Hashing
- Quantization
- Neighborhood graphs
- Most strategies address the curse of dimensionality problem

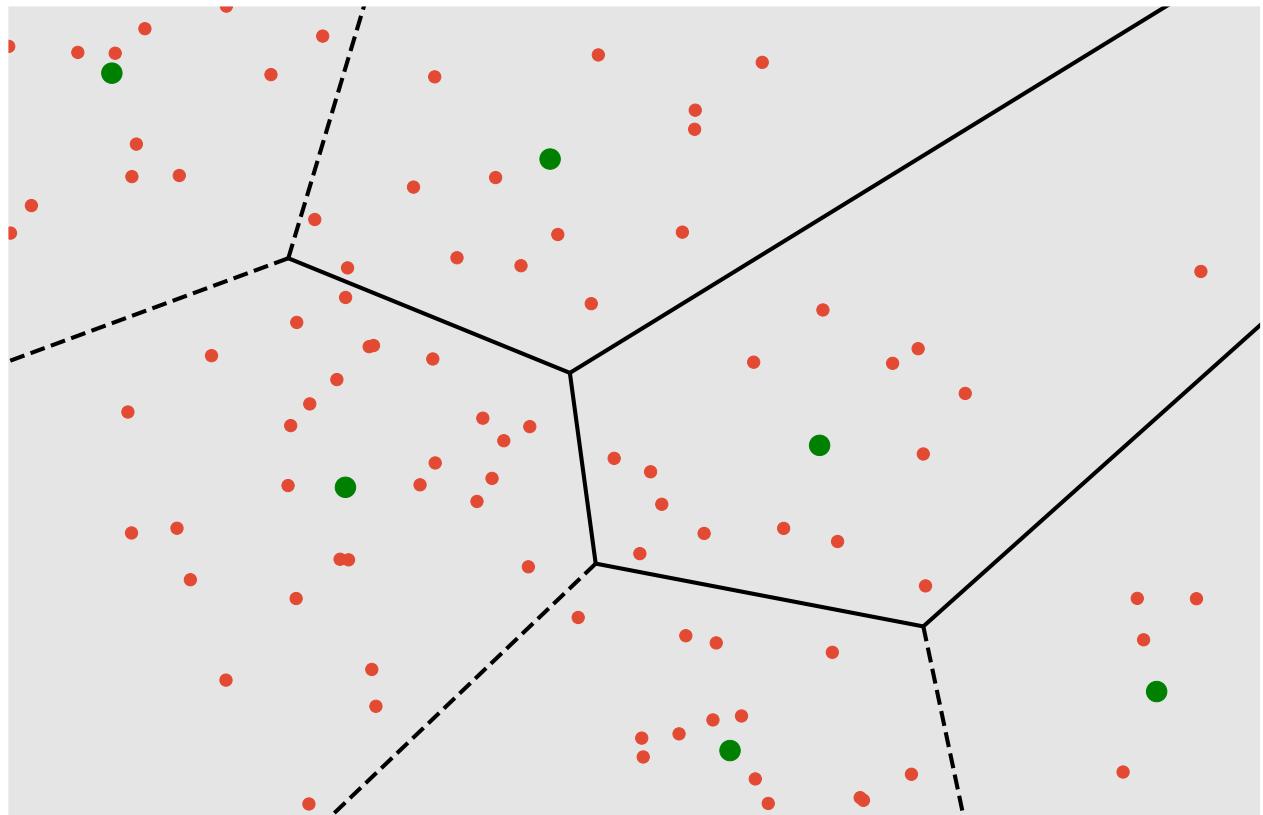
Quantization (I)

From n points to
 k centroids

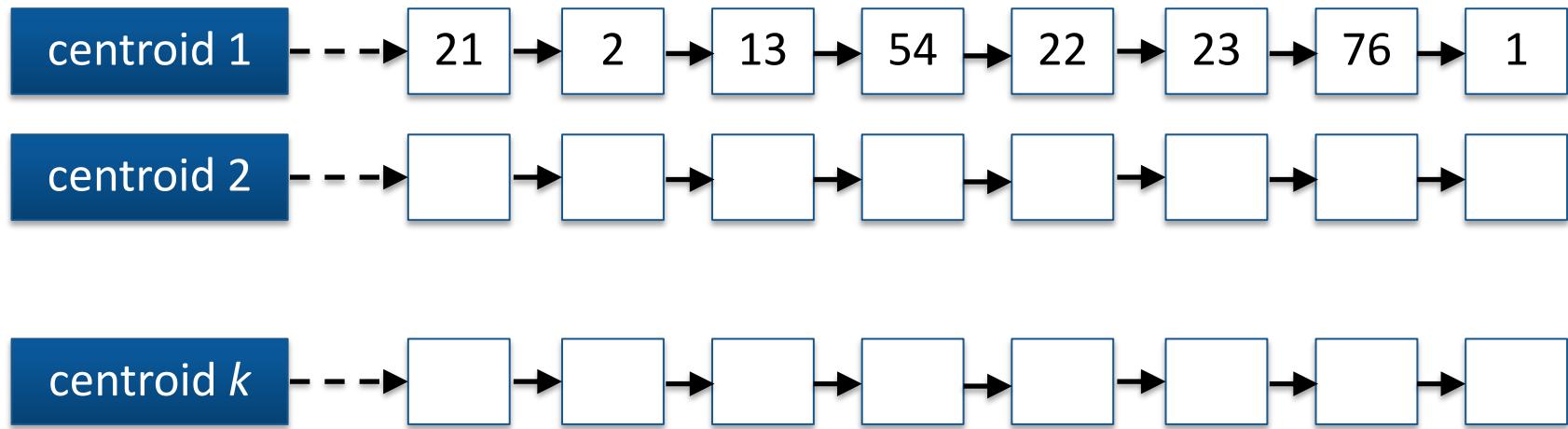


Quantization (II)

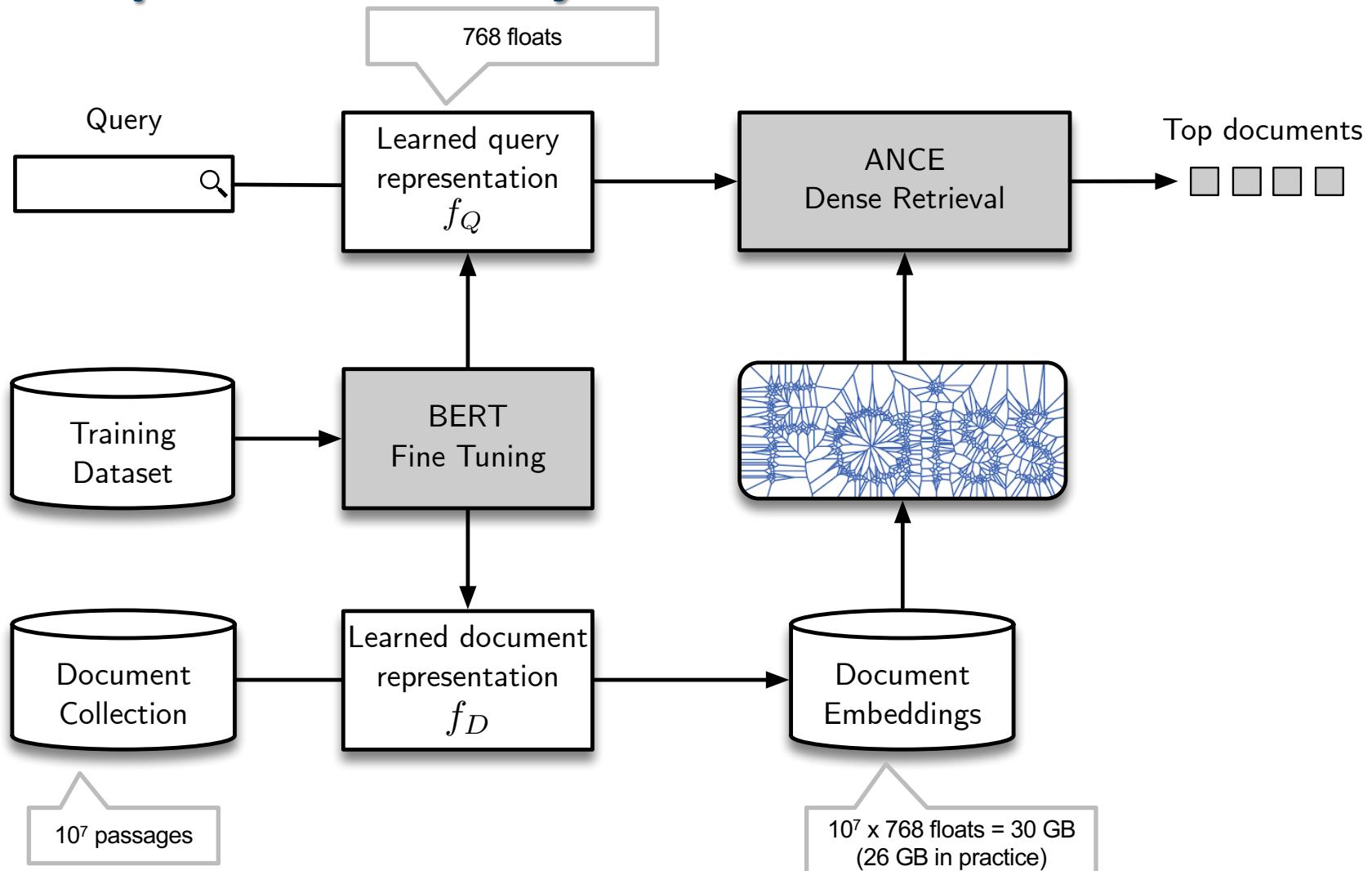
Use k-Means to select
centroids



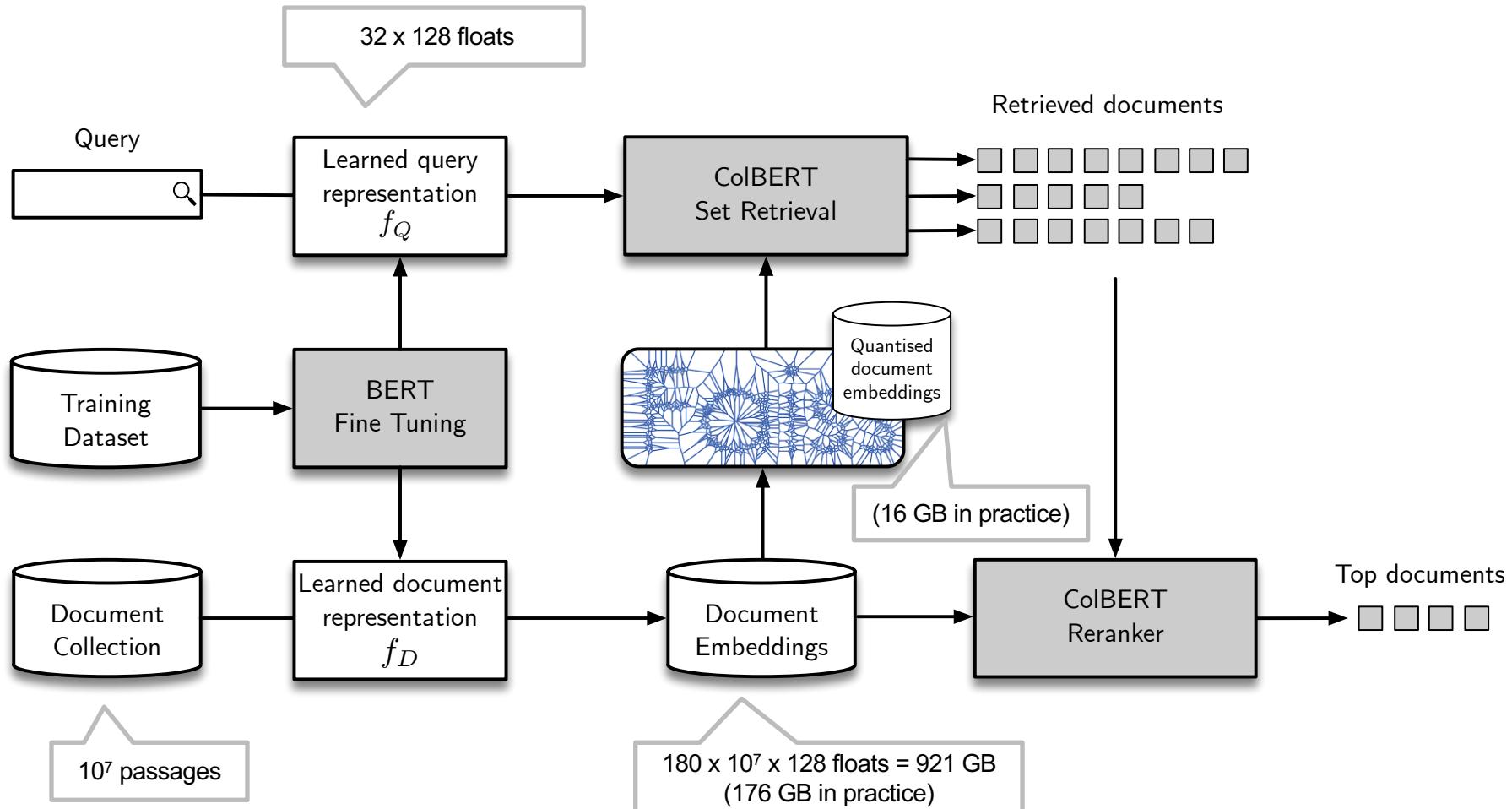
Quantization (III)



Dense Retrieval for Multiple Representation Systems



Approximate NN Dense Retrieval (ColBERT)



- ColBERT's token embeddings can be used to
 - select representative embeddings in a set of feedback documents using clustering
 - select discriminative embeddings among the representative embeddings using corresponding token's IDF
 - select the top tokens corresponding to discriminative embeddings for query expansions
- This approach works for both ranking and re-ranking scenarios

```
dense_e2e = pytcolbert.set_retrieve() >> pytcolbert.index_scorer(query_encoded=True, add_ranks=True)
if rerank:
    prf_pipe = (
        dense_e2e
        >> ColbertPRF(pytcolbert, k=k, fb_docs=fb_docs, fb_embs=fb_embs, beta=beta, return_docs=True)
        >> (pytcolbert.index_scorer(query_encoded=True, add_ranks=True) %1000)
    )
else:
    prf_pipe = (
        dense_e2e
        >> ColbertPRF(pytcolbert, k=k, fb_docs=fb_docs, fb_embs=fb_embs, beta=beta, return_docs=False)
        >> pytcolbert.set_retrieve(query_encoded=True)
        >> (pytcolbert.index_scorer(query_encoded=True, add_ranks=True) % 1000)
    )
return prf_pipe
```

ColBERT Optimisations

- **Single (ANCE) vs Multiple (ColBERT) Representations**
 - ANCE is more efficient than ColBERT
 - ColBERT is more effective than ANCE for MAP and MRR@10
 - ColBERT obtains better improvements than ANCE for queries that are the hardest for BM25, as well as for definitional queries, and those with complex information needs

Macdonald, Tonelotto, Ounis. On Single and Multiple Representations in Dense Passage Retrieval. IIR 2021

- **Embedding Pruning**
 - ColBERT does not need all 32 query embeddings, just 3
 - Same effectiveness, 2.65x speedup in efficiency

Tonelotto & Macdonald. Query Embedding Pruning for Dense Retrieval. CIKM 2021

- **Approximate Scoring**
 - ColBERT can use approximate scores from ANN to decrease the number of candidate documents being fully scored, e.g. 200 documents only
 - Same effectiveness, 2x speedup in efficiency

Macdonald & Tonelotto. On Approximate Nearest Neighbour Selection for Multi-Stage Dense Retrieval. CIKM 2021

From passages to documents



ANCE and ColBERT are designed for passages

- e.g. BERT has limited number of tokens for inference

For long documents, we need to break into passages

PyTerrier's **indexing** pipelines (c.f. part 2) offer a solution

```
import pyterrier_ance

ance_indexer = pt.text.sliding() >> pyterrier_ance.ANCEIndexer(
    checkpoint_path="../ance_model_checkpoint",
    index_path="/content/anceindex",
    num_docs=192509,
    text_attr='abstract' # COVID
)

ance_indexer.index(dataset.get_corpus_iter())
```

```
from pyterrier_ance import ANCERetrieval

ance_retriever = ANCERetrieval.from_dataset('trec-covid', 'ance_msmarco_psg') >> pt.text.max_passage()
```

ANCE Example

- Indexing

```
index = pt.datasets.get_dataset('trec-covid').get_index('terrier_stemmed')
ance_index = pt.datasets.get_dataset('trec-covid').get_index('ance_msMarco_psg')
```

- Retrieval

```
bm25_retriever = pt.BatchRetrieve(index, wmodel="BM25")
ance_retriever = ANCERetrieval.from_dataset('trec-covid', 'ance_msMarco_psg') >> pt.text.max_passage()
```

- Experiment

```
pt.Experiment(
    [bm25_retriever % 10, ance_retriever % 10],
    topics,
    qrels,
    eval_metrics=["map", "recip_rank", "P_10", "ndcg_cut_10", "mrt"],
    names=['BM25', 'ANCE'],
)
```

ColBERT Example

- End-to-end Retrieval

```
from pyterrier_colbert.ranking import ColBERTFactory

bm25_terrier_stemmed = pt.BatchRetrieve.from_dataset('vaswani', 'terrier_stemmed', wmodel='BM25')

factory = ColBERTFactory.from_dataset('vaswani', 'colbert_uog44k')
colbert_e2e = factory.end_to_end()
```

- Experiment

```
pt.Experiment(
    [bm25_terrier_stemmed % 10, colbert_e2e % 10],
    topics,
    qrels,
    eval_metrics=["map", "recip_rank", "P_10", "ndcg_cut_10", "mrt"],
    names=['BM25', 'ColBERT'],
)
```

Round Up

- **Index augmentation** can bring benefits from deep learning to traditional indices
 - Emphasize important terms, predict other terms that could match
- We can **use neural models** to directly learn scalar values to score terms in documents and store them in an inverted index
- We can use **dense retrieval systems** to **improve effectiveness** while **reducing the processing time** of queries
 - Represent **queries and documents** as one or more **embeddings** (single vs. multiple representation)
 - Embeddings are stored in a system providing **efficient support for NN search**
 - Depending on the size and number of embeddings, NN search can be **exact** or **approximate**

PyTerrier Github References



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- Doc2Query plugin: https://github.com/terrierteam/pyterrier_doc2query
- DeepCT plugin: https://github.com/terrierteam/pyterrier_deepct
- DeepImpact plugin: https://github.com/terrierteam/pyterrier_deepimpact
- SPLADE plugin: https://github.com/cmacdonald/pyt_splade
- ANCE plugin: https://github.com/terrierteam/pyterrier_ance
- ColBERT plugin: https://github.com/terrierteam/pyterrier_colbert

Questions

Practical Time



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The tutorial Github repo has links to the notebook for Part 3

- <https://github.com/terrier-org/searchsolutions2022-tutorial>
- Press the  Open in Colab link for each notebook to start a Colab session

Timings

- Practical: 15:45–16:15 GMT

Acknowledgements



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