



University  
of Glasgow



CIKM  
2021  
1-5 NOVEMBER

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

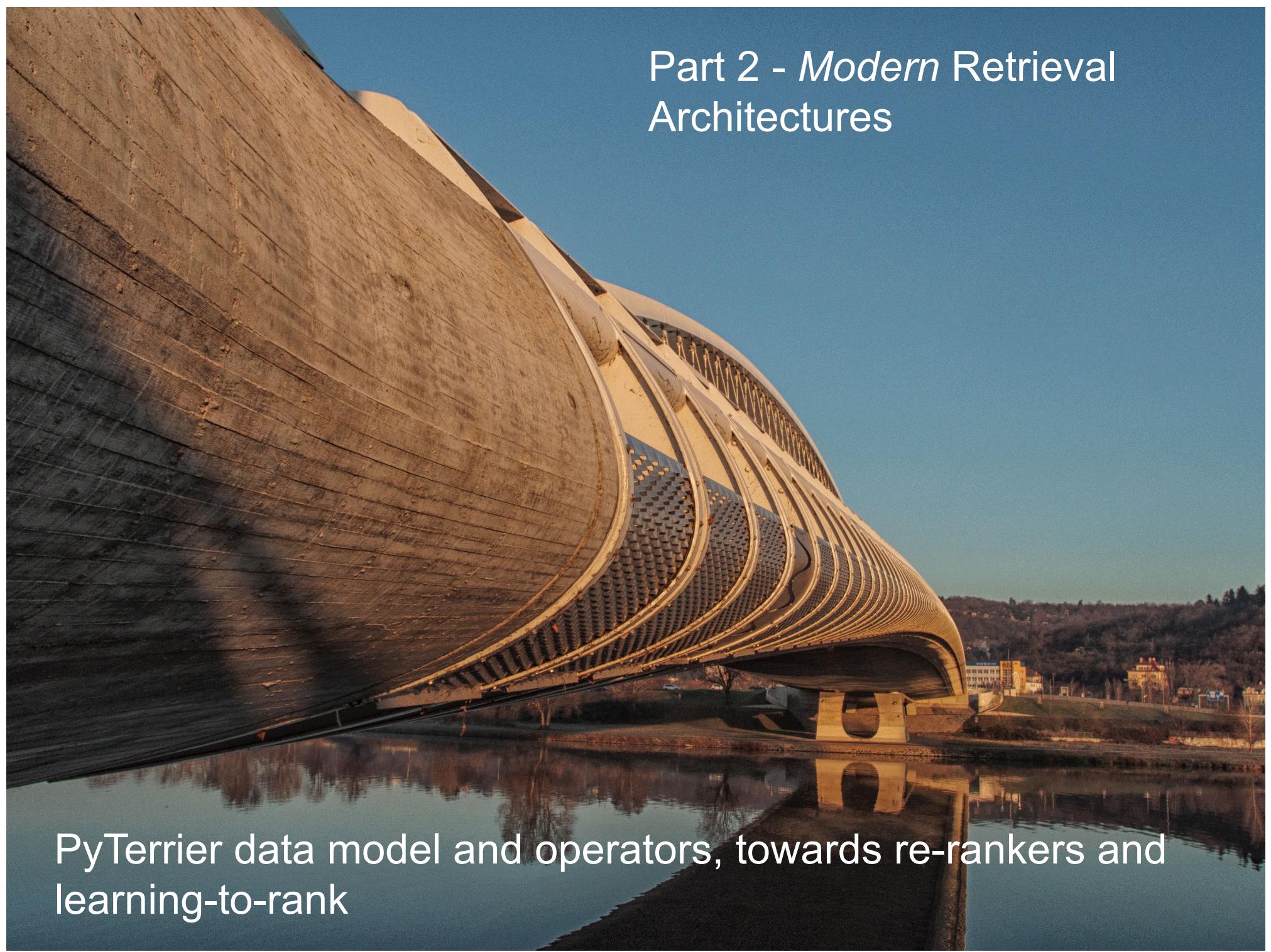
A CIKM 2021 tutorial with  
PyTerrier and OpenNIR

Sean MacAvaney\*  
Craig Macdonald\*  
Nicola Tonello\*

*Part 2 Starts at 1100 GMT (for Run 1)*

(\*Alphabetical ordering)

## Part 2 - *Modern Retrieval Architectures*



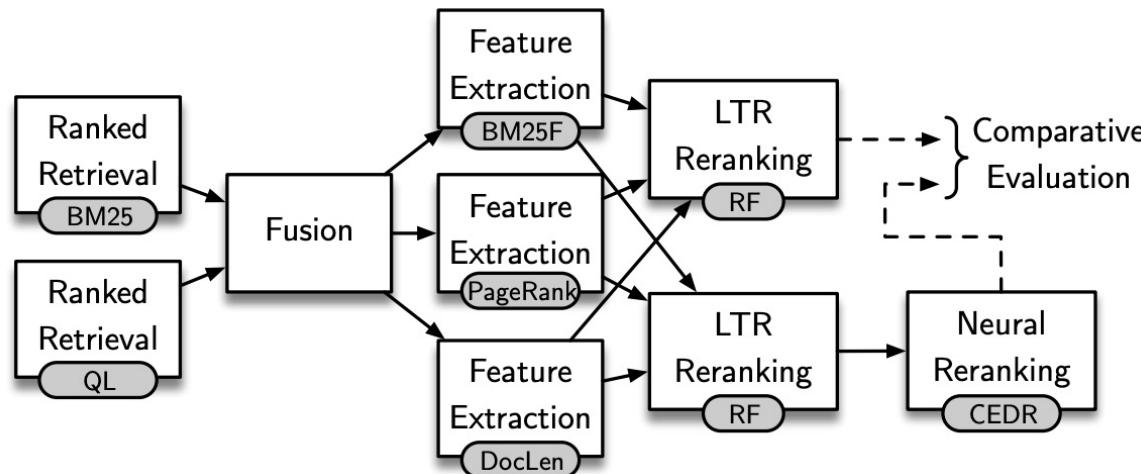
PyTerrier data model and operators, towards re-rankers and learning-to-rank

# Bridging to Modern Retrieval Architectures

In Part 1, we saw how to perform an experiment in a declarative manner, and perform simple retrievals

Actually, in IR, we have many retrieval and re-ranking components

Working with and combining result files to allow comparing complex ranking *pipelines* is tricky:



In Part 2, we generalise the architecture that has been seen into the PyTerrier data model and operators, to make it easy to build complex ranking pipelines

# *Modern Ranking Pipelines*



By using PyTerrier transformers and operators, we show that can easily express complex ranking pipelines in a **conceptual & declarative way**

Furthermore, we review **modern techniques** that fit into such ranking pipelines – such as...

- Query rewriting & pseudo-relevance feedback
- Learning to Rank

... and demonstrate how they can be implemented and extended within the PyTerrier framework

Finally, we also discuss the usefulness of PyTerrier for teaching IR

## *ILOs for this Session*



- ILO 2A. Understand modern retrieval architectures, such as re-rankers and ranking pipelines incorporating learning-to-rank strategies.
- ILO 2B. Understand how different ranking and re-ranking operations can be composed to make more complex ranking models, in a declarative manner and leveraging a specific data model.
- ILO 2C. Perform retrieval experiments within a Python notebook, including use of different features and learning-to-rank.
- ILO 2D. Understand the benefits of PyTerrier's declarative nature for teaching IR

# *Outline of Part 2*



**Part 2A: PyTerrier data model & transformers**

**Part 2B: Learning to Rank**

**Part 2C: Usefulness of PyTerrier for Teaching IR**

**Part 2D: Wrap up and move to practical session**



University  
of Glasgow

UNIVERSITÀ DI PISA

CIKM  
2021  
1-5 NOVEMBER

Part 2A

# PYTERRIER DATA MODEL AND TRANSFORMERS

# **PyTerrier makes it easy to construct complex IR pipelines**

**Key components:**

→ **A common data model**

- **A rich library of transformation functions**  
including classical retrieval, dense retrieval, LTR, neural re-ranking
- **Operators to combine transformers**

# Common Data Model



| <u>qid</u> | query                              |
|------------|------------------------------------|
| 0          | gold coast temperature november    |
| 1          | one way flight to gold coast price |
| ...        | ...                                |



| <u>docno</u> | text                           | title              |
|--------------|--------------------------------|--------------------|
| 0            | Science & Mathematics Physi... | The hot glowing... |
| 1            | School-Age Kids Growth &...    | Developmental...   |
| ...          | ...                            | ...                |



| <u>qid</u> | query                           | <u>docno</u> | <u>score</u> |
|------------|---------------------------------|--------------|--------------|
| 0          | gold coast temperature november | 15213        | 0.5134       |
| 0          | gold coast temperature november | 42635        | 0.3742       |
| 0          | gold coast temperature november | 26340        | 0.3223       |
| ...        | ...                             | ...          | ...          |



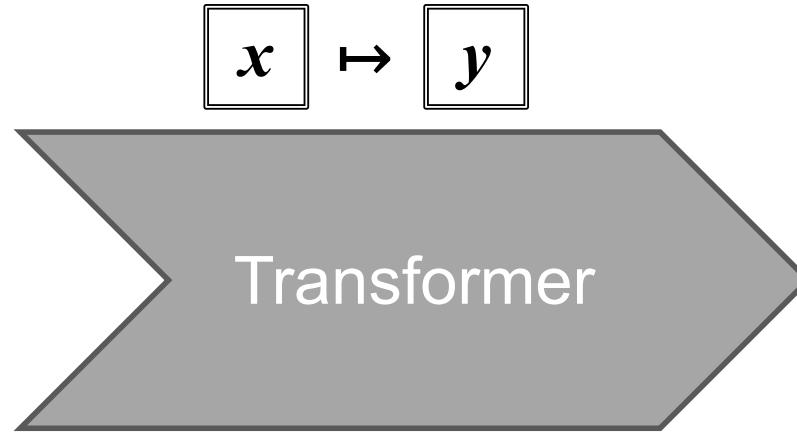
| <u>qid</u> | <u>docno</u> | <u>label</u> |
|------------|--------------|--------------|
| 0          | 15213        | 1            |
| ...        | ...          | ...          |

$RA \subset Q \times D$

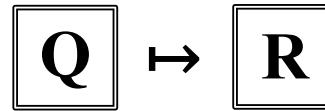
*(All of these are Pandas dataframes, i.e. relations)*

$R \subset Q \times D$

# Transformation Function Objects (Transformers)



# Transformation Function Objects (Transformers)

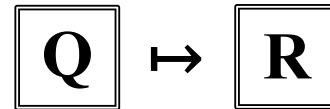


| qid | query                     |
|-----|---------------------------|
| 0   | gold coast temperature... |
| 1   | one way flight to gold... |
| ... | ...                       |



| qid | query         | docno | score  |
|-----|---------------|-------|--------|
| 0   | gold coast... | 15213 | 0.5134 |
| 0   | gold coast... | 42635 | 0.3742 |
| 0   | gold coast... | 26340 | 0.3223 |
| ... | ...           | ...   | ...    |

# Transformation Function Objects (Transformers)



| qid | query                     |
|-----|---------------------------|
| 0   | gold coast temperature... |
| 1   | one way flight to gold... |
| ... | ...                       |

Your latest  
retrieval  
approach...

| qid | query         | docno | score  |
|-----|---------------|-------|--------|
| 0   | gold coast... | 43242 | 0.1252 |
| 0   | gold coast... | 13746 | 0.1196 |
| 0   | gold coast... | 86454 | 0.0934 |
| ... | ...           | ...   | ...    |

# Transformation Function Objects (Transformers)



UNIVERSITÀ DI PISA



University  
of Glasgow

| Transformer Class       | Examples                              |
|-------------------------|---------------------------------------|
| $Q \mapsto R$ Retrieval | BM25, ColBERT, ANCE (dense retrieval) |

# Transformation Function Objects (Transformers)



UNIVERSITÀ DI PISA



University  
of Glasgow

| Transformer Class                 | Examples                              |
|-----------------------------------|---------------------------------------|
| $Q \rightarrow R$ Retrieval       | BM25, ColBERT, ANCE (dense retrieval) |
| $R \rightarrow Q$ Query Expansion | RM3, Bo1, etc.                        |

# Transformation Function Objects (Transformers)



| Transformer Class | Examples   |
|-------------------|--|
| $Q \rightarrow R$ | Retrieval<br>BM25, ColBERT, ANCE (dense retrieval) |
| $R \rightarrow Q$ | Query Expansion<br>RM3, Bo1, etc.                  |
| $R \rightarrow R$ | Re-ranking<br>Vanilla BERT, monoT5, etc.           |
| $Q \rightarrow Q$ | Query Re-writing<br>SDM, IntenT5, etc.             |

```
[12] index = pt.IndexRef.of('./cord19-index')
    bm25 = pt.TerrierRetrieve(index, wmodel='BM25')

    bm25(topics)
```

|       | <b>qid</b> | <b>docid</b> | <b>docno</b> | <b>rank</b> | <b>score</b> | <b>query</b>                                      |
|-------|------------|--------------|--------------|-------------|--------------|---|
| 0     | 1          | 83032        | dv9m19yk     | 0           | 11.865104    | what is the origin of covid 19                    |
| 1     | 1          | 109719       | kgifmjvb     | 1           | 11.412908    | what is the origin of covid 19                    |
| 2     | 1          | 135553       | wmfcey6f     | 2           | 11.397281    | what is the origin of covid 19                    |
| 3     | 1          | 68472        | 4dtk1kyh     | 3           | 10.805808    | what is the origin of covid 19                    |
| 4     | 1          | 114244       | cniyembt     | 4           | 10.763978    | what is the origin of covid 19                    |
| ...   | ...        | ...          | ...          | ...         | ...          | ...   |
| 49995 | 50         | 146581       | 2o8u4eny     | 995         | 17.073736    | what is known about an mrna vaccine for the sa... |
| 49996 | 50         | 76091        | fayr38c7     | 996         | 17.071367    | what is known about an mrna vaccine for the sa... |
| 49997 | 50         | 102986       | jlrk8fg0     | 997         | 17.070633    | what is known about an mrna vaccine for the sa... |
| 49998 | 50         | 183502       | imwa033f     | 998         | 17.070633    | what is known about an mrna vaccine for the sa... |
| 49999 | 50         | 95511        | j5j3hxxw     | 999         | 17.070486    | what is known about an mrna vaccine for the sa... |

## BM25 Results

```
[22] rm3 = pt.rewrite.RM3(index)
      rm3(bm25(topics))
```

|   | qid | query   | query_0   |
|---|-----|---|---|
| 0 | 1   | 19^0.200000018 origin^0.325867474 covid^0.2000... | what is the origin of covid 19                    |
| 1 | 10  | 19^0.085714296 across^0.019867204 epidem^0.029... | has social distancing had an impact on slowing... |
| 2 | 11  | walk^0.030252097 urolog^0.042577028 optim^0.04... | what are the guidelines for triaging patients ... |
| 3 | 12  | older^0.041744966 best^0.100000009 quarantin^0... | what are best practices in hospitals and at ho... |
| 4 | 13  | contact^0.026820807 possibl^0.021502888 outbre... | what are the transmission routes of coronavirus   |
| 5 | 14  | centuri^0.020000001 manag^0.020000001 contain^... | what evidence is there related to covid 19 sup... |
| 6 | 15  | bodi^0.167176828 can^0.120000005 cold^0.034101... | how long can the coronavirus live outside the ... |
| 7 | 16  | stabl^0.152924821 dai^0.042988449 surfac^0.218... | how long does coronavirus remain stable on sur... |

Q

Expanded query

Original query

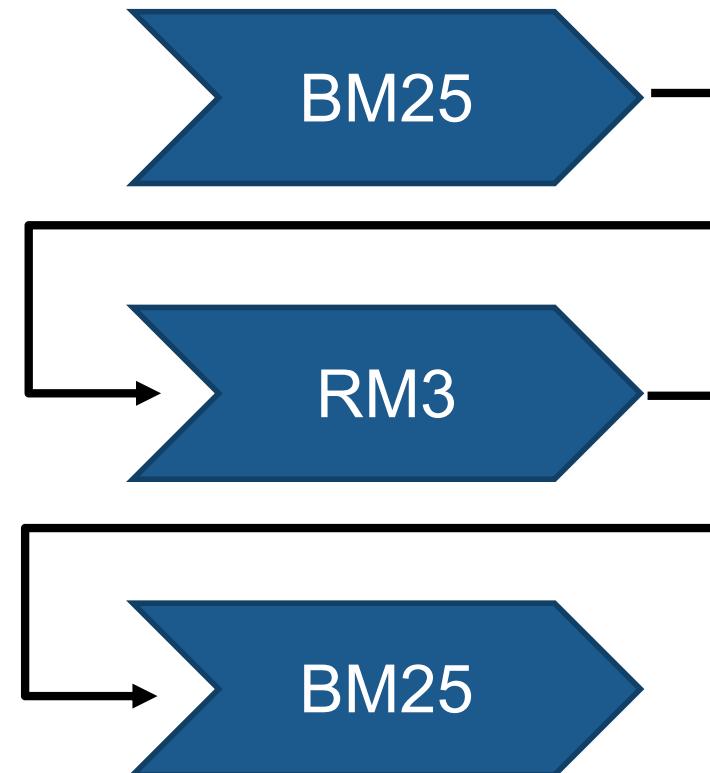
[23] `bm25(rm3(bm25(topics)))`

|       | <b>qid</b> | <b>docid</b> | <b>docno</b> | <b>rank</b> | <b>score</b> |   | <b>query</b>                     | <b>query_0</b> |
|-------|------------|--------------|--------------|-------------|--------------|---|----------------------------------|----------------|
| 0     | 1          | 109719       | kgifmjvb     | 0           | 19.316403    | 19^0.200000018 origin^0.325867474 covid^0.2000... | what is the origin of covid 19   |                |
| 1     | 1          | 135553       | wmfcey6f     | 1           | 19.281917    | 19^0.200000018 origin^0.325867474 covid^0.2000... | what is the origin of covid 19   |                |
| 2     | 1          | 83032        | dv9m19yk     | 2           | 11.101417    | 19^0.200000018 origin^0.325867474 covid^0.2000... | what is the origin of covid 19   |                |
| 3     | 1          | 103317       | ec8lpgl3     | 3           | 11.044733    | 19^0.200000018 origin^0.325867474 covid^0.2000... | what is the origin of covid 19   |                |
| 4     | 1          | 114244       | cniyembt     | 4           | 10.865901    | 19^0.200000018 origin^0.325867474 covid^0.2000... | what is the origin of covid 19   |                |
| ...   | ...        | ...          | ...          | ...         | ...          | ...   | ...                              | ...            |
| 49995 | 9          | 164232       | 6djfpvz7     | 995         | 6.987903     | 19^0.150000006 covid^0.150000006 mp^0.05102380... | how has covid 19 affected canada |                |
| 49996 | 9          | 68116        | 6k8vedy3     | 996         | 6.983706     | 19^0.150000006 covid^0.150000006 mp^0.05102380... | how has covid 19 affected canada |                |
| 49997 | 9          | 132455       | 9z0azq8y     | 997         | 6.983706     | 19^0.150000006 covid^0.150000006 mp^0.05102380... | how has covid 19 affected canada |                |
| 49998 | 9          | 132456       | wfss2j7v     | 998         | 6.983706     | 19^0.150000006 covid^0.150000006 mp^0.05102380... | how has covid 19 affected canada |                |
| 49999 | 9          | 163423       | n76892ur     | 999         | 6.981468     | 19^0.150000006 covid^0.150000006 mp^0.05102380... | how has covid 19 affected canada |                |

## Results of BM25 over RM3

# Operators

bm25 ( rm3 (bm25 (topics) ) )

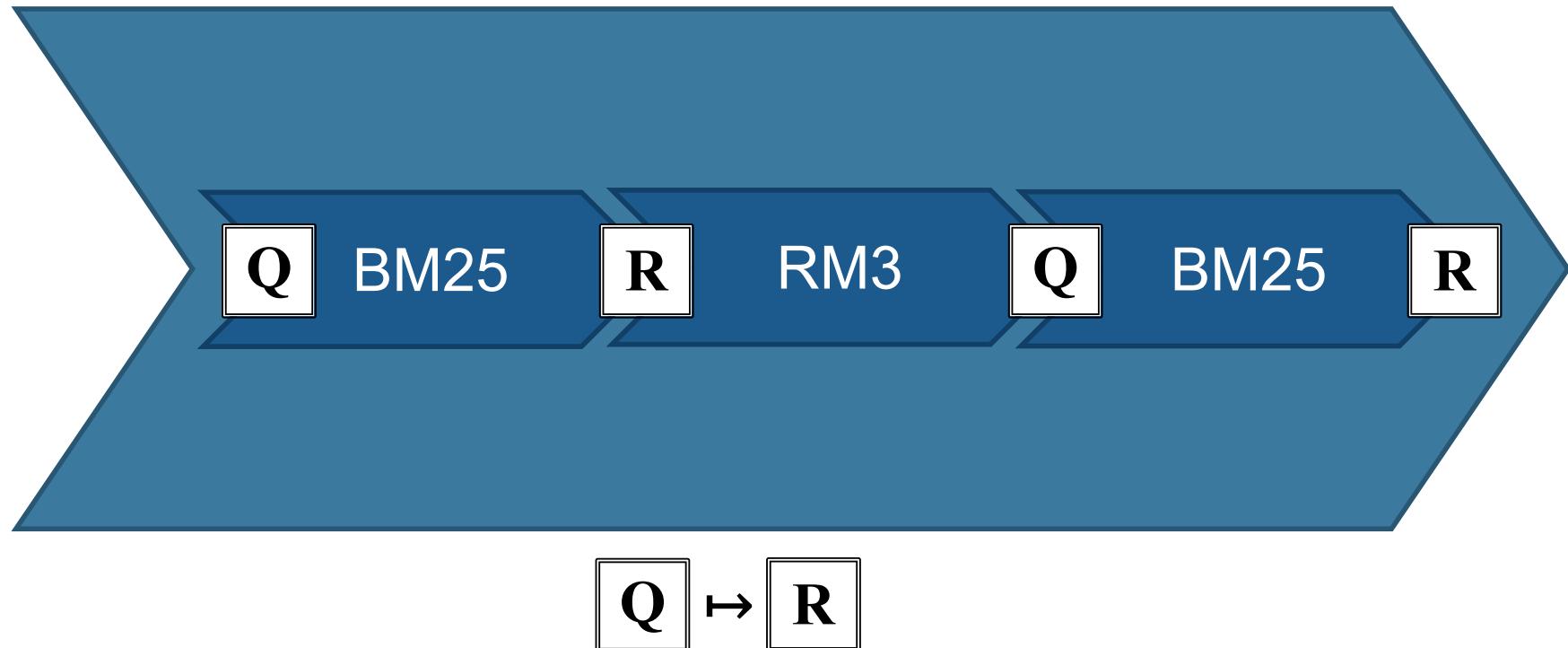


# Operators

A better way: the “then” operator

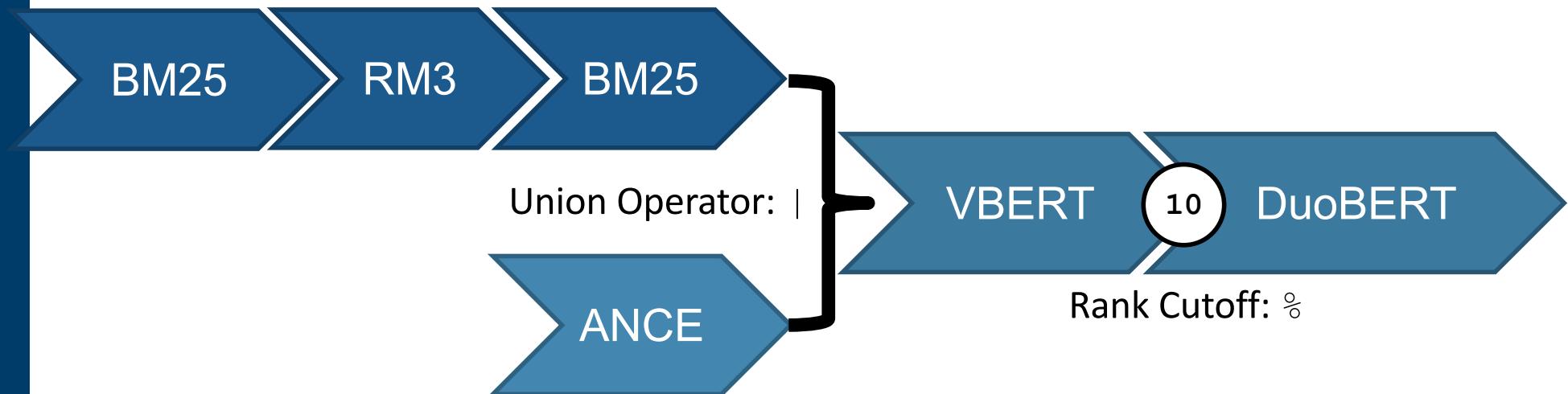
A >> B (A then B)

pipeline = bm25 >> rm3 >> bm25



# Operators

```
pipeline = bm25 >> rm3 >> bm25
```



Other operators available, including caching, linear score combination, etc.

# Transformer Operators



## PyTerrier has a suite of operators on transformers

| Op. | Name                          | Description  |
|-----|-------------------------------|--|
| >>  | <i>then<br/>(composition)</i> | Pass the output from one transformer to the next transformer                   |
| +   | <i>linear combine</i>         | Sum the query-document scores of the two retrieved results lists               |
| *   | <i>scalar product</i>         | Multiply the query-document scores of a retrieved results list by a scalar     |
| **  | <i>feature union</i>          | Combine two retrieved results lists as features                                |
|     | <i>set union</i>              | Make the set union of documents from the two retrieved results lists           |
| &   | <i>set intersection</i>       | Make the set intersection of the two retrieved results lists                   |
| %   | <i>rank cutoff</i>            | Shorten a retrieved results list to the first $K$ elements                     |
| ^   | <i>concatenate</i>            | Add the retrieved results list from one transformer to the bottom of the other |

Our ICTIR 2020 paper on PyTerrier provides relational algebra semantics for each operator

- Let's look at a few others...

# Lets look at some retrieval use cases

## 1. Linear Combinations

```
linear = 2 * pt.BatchRetrieve(index, wmodel="BM25")  
+ 3 * pt.BatchRetrieve(index, wmodel="DPH")
```

Final Output Type: R

Output Types: R

## 2. Sequential Dependence Model (proximity)

```
sdm = pt.rewrite.SequentialDependence() >> bm25
```

Input Type: Q

| qid | query                |
|-----|----------------------|
| 0   | 1 chemical reactions |

Output Type: Q

Output Type: R

| qid | query   |
|-----|---|
| 0   | 1<br>#combine:0=0.1:wmodel=org.terrier.matching.models.dependence.pBiL(#1(chemical reactions))<br>#combine:0=0.1:wmodel=org.terrier.matching.models.dependence.pBiL(#uw8(chemical reactions))<br>#combine:0=0.1:wmodel=org.terrier.matching.models.dependence.pBiL(#uw12(chemical reactions)) |

# Lets look at some retrieval use cases

## 1. Linear Combinations

```
linear = 2 * pt.BatchRetrieve(index, wmodel="BM25")  
       + 3 * pt.BatchRetrieve(index, wmodel="DPH")
```

Final Output Type: R

Output Types: R

## 2. Sequential Dependence Model (proximity)

```
sdm = pt.rewrite.SequentialDependence() >> bm25
```

Input Type: Q

Output Type: Q

Output Type: R

## 3. Pseudo-relevance feedback (QE)

```
rm3 = bm25 >> pt.rewrite.RM3(index) >> bm25
```

Output Type: R

Output Type: Q

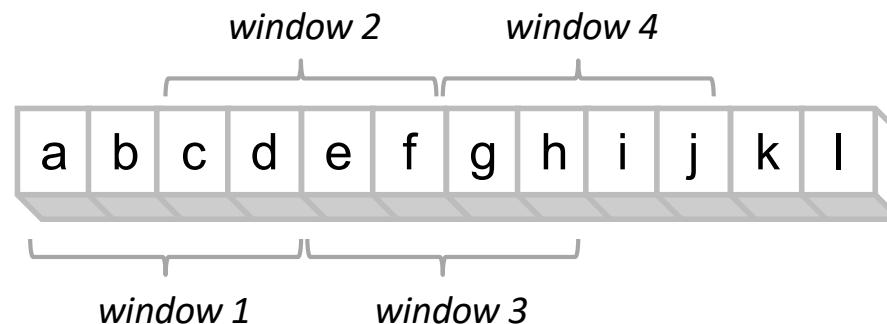
Output Type: R

# *Retrieval Use Case: Passaging*

Some long documents may only contain a relevant passage about the query

- In such cases, **passage-based retrieval** can benefit effectiveness
- It's also useful for neural models that do not scale efficiently to long documents

For instance, applying a **sliding window** over the text of the document to get passages, and scoring each passage individually



Window size 4  
Window stride 2

- Some settings benefit from prepending the title to each passage

# Retrieval Use Case: Passaging

Lets try this out:

```
Bm25maxp = pt.BatchRetrieve(index, wmodel="BM25", metadata=[“docno”, “text”])  
>> pt.text.sliding(“text”, length=150, stride=75)  
>> pt.text.scorer(wmodel="BM25")  
>> pt.text.max_passage()
```

| sliding(“text”, 2, 1) |       |         |  |
|-----------------------|-------|---------|--|
| BR output (R w/ text) |       | output  |  |
| qid                   | docno | text    |  |
| q1                    | d1    | a b c d |  |
|                       |       |         |  |
| q1                    | d1%p1 | a b     |  |
| q1                    | d1%p2 | b c     |  |
| q1                    | d1%p3 | c d     |  |

| textscorer output (R) |       |      |       |
|-----------------------|-------|------|-------|
| qid                   | docno | rank | score |
| q1                    | d1%p5 | 0    | 5.0   |
| q1                    | d2%p4 | 1    | 4.0   |
| q1                    | d1%p3 | 2    | 3.0   |
| q1                    | d1%p1 | 3    | 1.0   |

| max_passage output |       |      |       |
|--------------------|-------|------|-------|
| qid                | docno | rank | score |
| q1                 | d1    | 0    | 5.0   |
| q1                 | d2    | 1    | 4.0   |

(Some attributes omitted for brevity, e.g. query)

Of course, we could change `pt.text.scorer()` to be a more advanced approach, e.g. BERT-based neural re-ranker [Part 3]

# *Indexing Pipelines*



## Applying passaging during retrieval can be expensive

- Can it be applied at indexing time?
- There are also other techniques that we will see in Parts 3 & 4 (e.g. DeepCT, doc2query) that can be applied at indexing time

The `>>` operator can also be applied to create **indexing pipelines**

Hence, a MaxPassage retrieval can also be achieved as follows

```
indexer = pt.text.sliding("text", length=150, stride=75)
        >> pt.IterDictIndexer()
indexref = indexer.index(dataset.get_corpus_iter())
MaxP = pt.BatchRetrieve(indexref, wmodel="BM25")
        >> pt.text.max_passage()
```

Note that while transformers operate on a dataframes, we might not fit an entire corpus into a dataframe

- Hence indexing pipelines perform transparent batching of documents

# Transformer Implementations (1/2)



UNIVERSITÀ DI PISA



All transformers implement a base class called `TransformerBase`

- This has all the methods to support operator overloading, e.g. `__add__()`, `__rshift__()`

However, creating your own transformer is easy using `pt.apply` functions. These create a new transformer from your own function

- For instance, to rewrite a query – `pt.apply.query()`

```
# this will remove pre-defined stopwords from the query
stops=set(["and", "the"])

# a naieve function to remove stopwords
def _remove_stops(q):
    terms = q["query"].split(" ")
    terms = [t for t in terms if not t in stops]
    return " ".join(terms)

# a query rewriting transformer applying _remove_stops
p1 = pt.apply.query(_remove_stops) >> pt.BatchRetrieve(index, wmodel="DPH")
```

- `pt.apply.query()` returns a transformer, which applies `_remove_stops` to each query. The previous query is saved in the `query_0` attribute

# *Transformer Implementations (2/2): pt.apply.doc\_score()*



`pt.apply.doc_score()` uses a function that returns a score

```
# this transformer will subtract the number of character in the query
# from the score of each retrieved document

p = pt.BatchRetrieve(index, wmodel="DPH") >>
    pt.apply.doc_score(lambda doc : doc["score"] - len(doc["query"]))
```

In this way, ANY function that can return a number can be used for retrieval

- i.e. any (SOTA) approach that can be expressed as a function of the **text of the document** and the **text of the query** can be used as a re-ranker in PyTerrier

`pt.apply.doc_score()` & `pt.apply.query()` have known semantics

- There are also advanced “generic” apply functions that can apply an arbitrary change of a dataframe as a transformer

# ***Summary: PyTerrier Standard Transformers***



## **Retrieval**

- `pt.BatchRetrieve()`

## **Query Rewriting/Expansion**

- `pt.rewrite.SequentialDependence()`
- `pt.rewrite.RM3()`
- `pt.rewrite.Bo1QueryExpansion()`
- `pt.rewrite.KLQueryExpansion()`

## **Text**

- `pt.text.get_text()`
- `pt.text.scorer()`
- `pt.text.sliding()`
- `pt.text.max_passage()` `first_passage()` `mean_passage()`  
`kmaxavg_passage()`

In Part 2B, we'll see learning-to-rank transformers

In Parts 3 & 4, we'll see more add-on transformers for neural re-ranking and end-to-end

# **Summary: PyTerrier Utility Transformers**



UNIVERSITÀ DI PISA



## **Apply functions as pipelines**

- `pt.apply.doc_score(_fn)`
- `pt.apply.query(_fn)`
- `pt.apply.generic(_fn)`
- `pt.apply.by_query(_fn)`
- `pt.apply.<ANY COLUMN NAME>(_fn)`
- `pt.apply.<ANY COLUMN NAME>(drop=True)`
- `pt.apply.rename(columns={"old": "new"})`

User-defined functions to  
change the pipeline in  
arbitrary ways

## **Inspect within a transformer pipeline**

- `pt.debug.print_rows()`
- `pt.debug.print_columns()`
- `pt.debug.print_num_rows()`

# Summary: PyTerrier Transformer Operators



| Op. | Name                                | Description  |
|-----|-------------------------------------|--|
| >>  | <i>then</i><br><i>(composition)</i> | Pass the output from one transformer to the next transformer                   |
| +   | <i>linear combine</i>               | Sum the query-document scores of the two retrieved results lists               |
| *   | <i>scalar product</i>               | Multiply the query-document scores of a retrieved results list by a scalar     |
| **  | <i>feature union</i>                | Combine two retrieved results lists as features <b>(Part 2B)</b>               |
|     | <i>set union</i>                    | Make the set union of documents from the two retrieved results lists           |
| &   | <i>set intersection</i>             | Make the set intersection of the two retrieved results lists                   |
| %   | <i>rank cutoff</i>                  | Shorten a retrieved results list to the first $K$ elements                     |
| ^   | <i>concatenate</i>                  | Add the retrieved results list from one transformer to the bottom of the other |

# Reflections (1)



Transformers build on PyTerrier's extensible data model

- defined on IR-level objects: queries and documents
- One could imagine more data model columns for, e.g., ConvQA

Operators, defined with clear relational algebra semantics, allow different transformers to be easily combined

- The structure of the pipeline implementations are close to their conceptual design
- This is all declarative in nature - we aren't writing code that iterates over any queries or documents (no for loops 😊!)

New techniques can be used and evaluated within a unified framework

- `pt.apply.doc_score()` can be easily applied to integrate new state-of-the-art approaches

# Reflections (2)

Thinking in this manner allows many techniques to be easily instantiated... and new techniques to be designed.

- We have created pipelines for integrating PRF into dense retrieval
- Both of our CIKM papers making advances on ColBERT are designed and implemented as variants of pipelines

```
import pyterrier_colbert.pruning

#CIKM 2021 Approximate Scoring paper: only retrieve 200 candidates for exact re-ranking
ann_pipe = (factory.ann_retrieve_score() % 200) >> factory.index_scorer(query_encoded=True)

#CIKM 2021 query embeddings paper: only keep the 9 tokens with highest ICF
qep_pipe5 = (factory.query_encoder()
             >> pyterrier_colbert.pruning.query_embedding_pruning(factory, 5)
             >> factory.set_retrieve(query_encoded=True)
             >> factory.index_scorer(query_encoded=False)
         )
```

- qep\_pipe5 has the following transformations

$[qid, query] \quad \text{Q} \quad \gg \quad [qid, query, query\_embs] \quad \text{Q+} \quad \gg \quad [qid, query, query\_embs, docno] \quad \text{R+} \quad \gg \quad [qid, query, query\_embs, docno, rank, score] \quad \text{R+}$

Even if we have a single coherent end-to-end retrieval model, for deployment, we may need things like query-biased snippets, which are pipelined after retrieval...

# *Building your own*



We have been extending PyTerrier with “plugins” for various models - some you will see in Parts 3 & 4:

- pyt\_t5, pyt\_colbert, pyt\_ance, pyt\_doc2query, pyt\_deepimpact

Its easy to make a retrieval function into a PyTerrier transformer:

- Make a class that extends TransformerBase
- Define a transform() method that takes as input a Q dataframe and returns an R dataframe
- pt.apply functions can reduce the boilerplate even further

Indexers also can be supported

- Make a class that extends TransformerBase
- Define an index() method that takes iterable of dicts each containing a document.

Then its plain sailing 🌟 🌟 🌟 !

# *Reasoning on Transformer Pipelines (1/2)*



Validation

*(Coming Soon!)*

*pipe.validate()*

- Lots of transformers, lots of operators. Is my pipeline expression valid? Do I have the correct columns?
- We can know the desired input & actual output columns of each transformer
  - E.g. BatchRetrieve needs qid and query columns, sliding() needs text column etc
- Similarly, we know actual inputs and desired outputs of an entire pipeline
  - E.g. We know that queries have qid and query columns
  - E.g. We know that we need qid, docno, score to evaluate a ranking
- Therefore, it is possible to *validate* (type check) a pipeline before execution – check if each transformer stage emits the necessary columns for the next stage
  - $Q + Q$  would throw an error – we cannot make a linear combination of  $Q$
- This would be transparent to most users; Some “generic” `pt.apply` operations would need to specify their input and output columns to ensure validation

# *Reasoning on Transformer Pipelines (2/2)*

## Optimisation      *pipe = pipe.compile()*

- Pipelines with identical semantics can be written in different ways  
`br1 = pt.BatchRetrieve(index) % 10`  
`br2 = pt.BatchRetrieve(index, num_results=10)`
- In our ICTIR 2020 paper, we showed that pipelines can be rewritten into equivalent but faster equivalent pipelines as a form of graph rewriting

## Parallelisation      *pipe = pipe.parallel(2)*

- Transformers can be parallelized using different processes, or even to different machines
- This is because the primary key attributes of each transformer are known, so we can partition a Q or R dataframe by qid; a D dataframe by docno

# Tuning: GridScan & GridSearch

Often, we have transformers with parameters, e.g.

```
BM25 = pt.BatchRetrieve(index, wmodel="BM25",
                         controls={"c" : 0.75, "bm25.k_1": 0.75, "bm25.k_3": 0.75})
```

Which we need to tune... pt.GridSearch and pt.GridScan

```
pt.GridSearch(
    BM25,
    {BM25: {
        "c" : [0, 0.2, 0.4, 0.6, 0.8, 1 ],
        "bm25.k_1": [ 0.3, 0.6, 0.9, 1.2, 1.4, 1.6, 2 ],
        "bm25.k_3": [ 0.5, 2, 4, 6, 8, 10, 12, 14, 20 ]
    }},
    train_topics,
    train_qrels,
    "map")
```

*Transformer pipeline to use*

*Dictionary mapping transformer to parameter to value range*

*Topics, qrels, measure*

- GridSearch – find the best setting; GridScan – evaluate all settings
- Also a KFoldGridSearch facilitates cross-validation settings



University  
of Glasgow

UNIVERSITÀ DI PISA

CIKM  
2021  
1-5 NOVEMBER

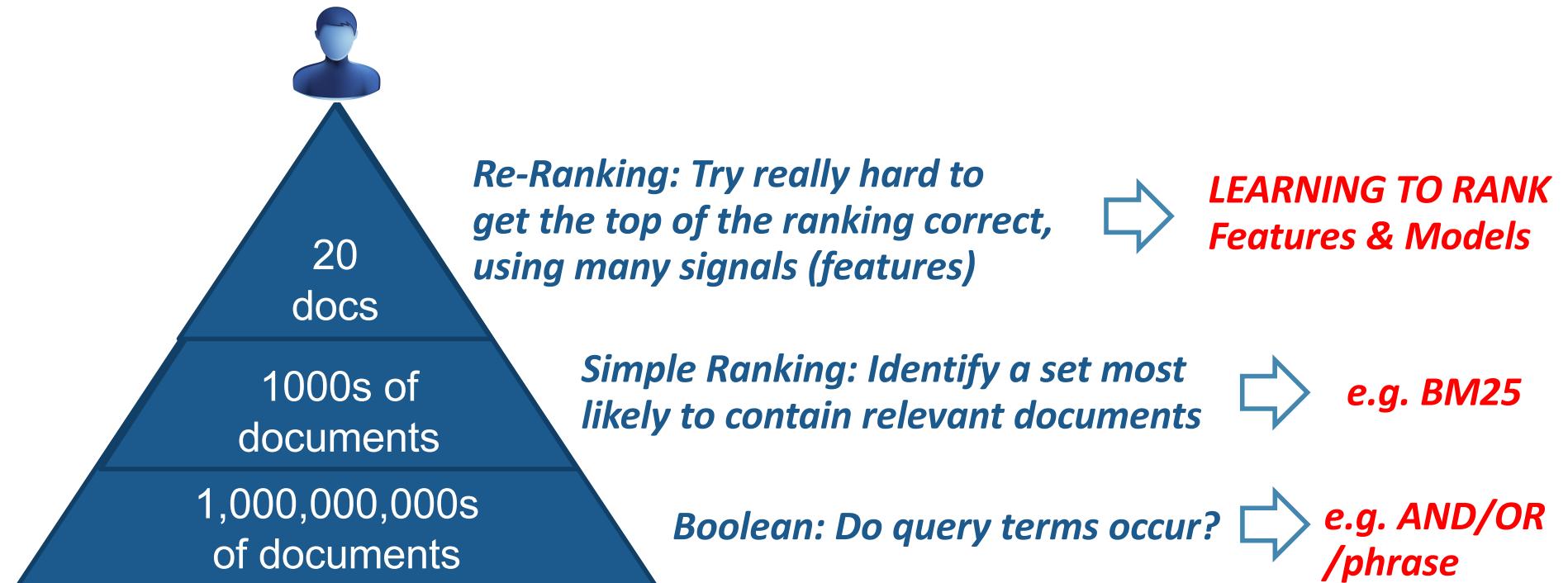
Part 2B

# LEARNING TO RANK

# Ranking Cascades

In many search scenarios, the ranking process can be seen as a series of cascades [1]

- Rank some documents
- Pass top-ranked onto next cascade for refined re-ranking



[1] J Pederson. Query understanding at Bing. SIGIR 2010 Industry Day.

## How to choose term weighting models?

- Term weighting models have different **assumptions** about how relevant documents should be retrieved

### Also:

- **Proximity-models:** close co-occurrences matter more
- **Priors:** documents with particular lengths or URL/inlink distributions matter more
- **Field-based models:** term occurrences in different fields matter differently (e.g. BM25F, PL2F)
- **Query Features:** Long queries, difficult queries, query type

How to combine all these easily and appropriately?

# **Learning to Rank definition**

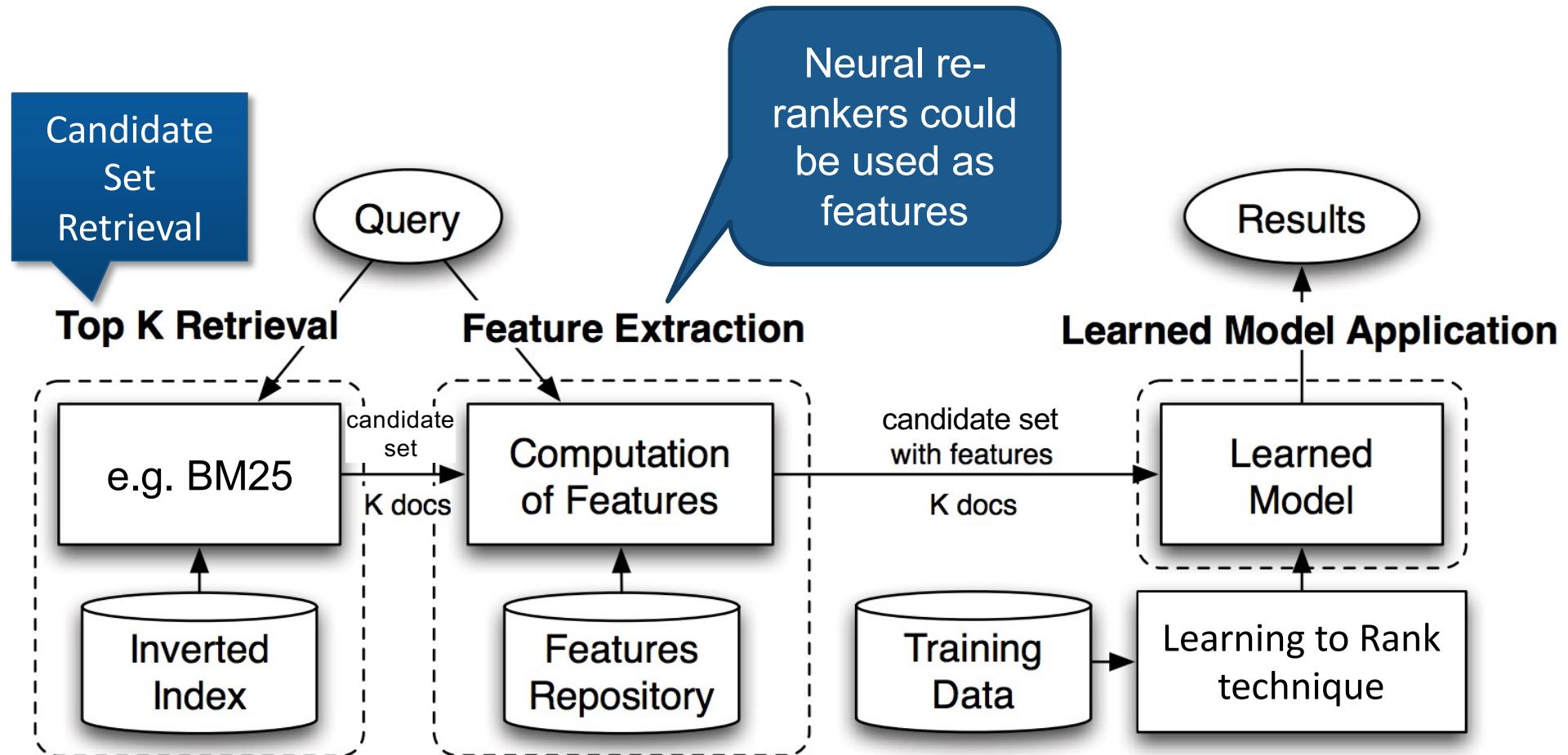


**Learning to rank describes the application of tailored machine learning techniques, applied to re-rank a candidate set of retrieved documents**

- This requires hand-engineered features (incl. weighting models) as relevance signals
- LTR techniques should focus on getting the top-ranked documents ranked correctly, using adapted loss functions:
  - Pointwise: e.g. regression – just tries to predict document label
  - Pairwise: tries to get pairs of documents with differing label correctly ranked
  - Listwise: tries to get most relevant document at top of ranking

**Different model forms are popular: linear, trees, neural nets**

# LTR, Schematically...



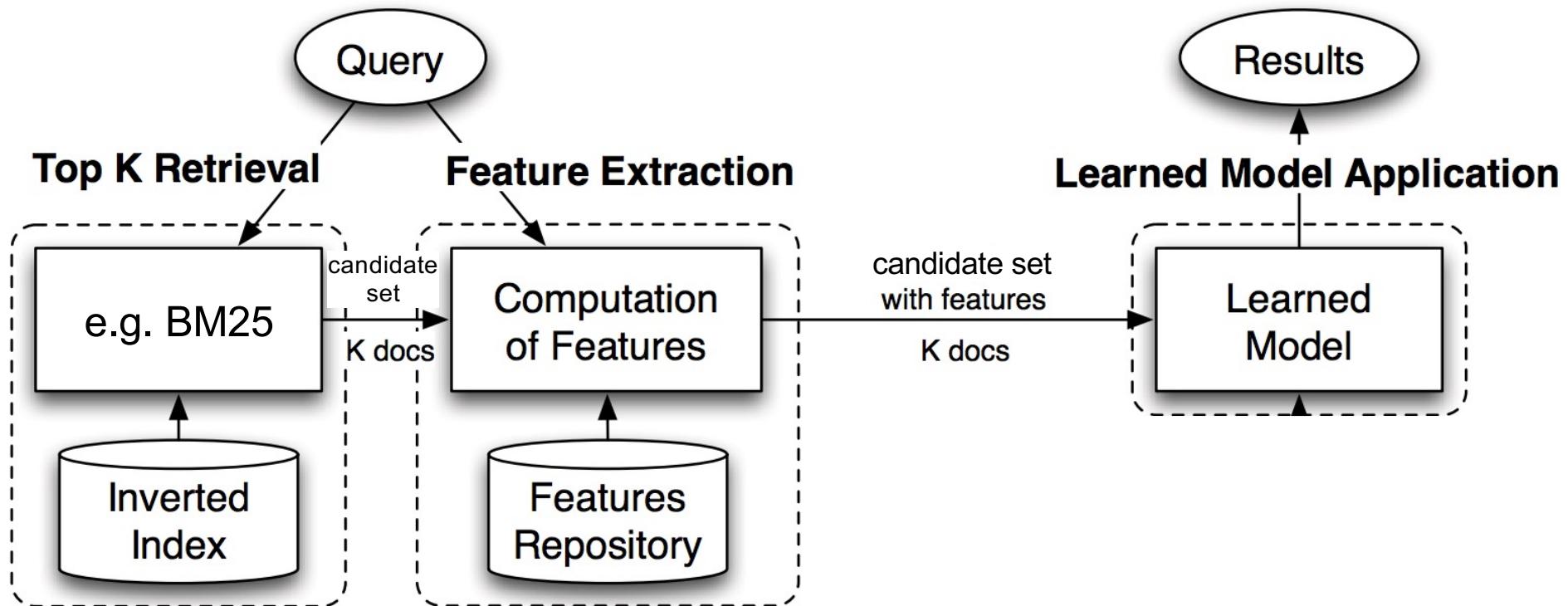
# Types of Features



Typically, commercial search engines use hundreds of features for ranking documents, usually categorised as follows:

| Name                       | Varies depending on... |          | Examples   |
|----------------------------|------------------------|----------|--|
|                            | Query                  | Document |  |
| Query Dependent Features   | ✓                      | ✓        | Weighting models, e.g. BM25, PL2<br>Proximity models, e.g. Markov Random Fields<br>Field-based weighting models, e.g. PL2F<br>Neural re-rankers [Part 3] |
| Query Independent Features | X                      | ✓        | PageRank, number of inlinks<br>Spamminess  |
| Query Features             | ✓                      | X        | Query length, Query performance predictors<br>Presence of entities   |

# Matching LTR concepts to PyTerrier...



In PyTerrier:

```
pt.BatchRetrieve()  
**  
pt.BatchRetrieve() % K >> pt.apply_doc_score() ** >> pt.ltr.apply_learned_model()  
::
```

# ***“Featured” Data Model***



**Retrieved documents with **features**,  $R_f$ :**

| <u>qid</u> | query | <u>docno</u> | score | rank | <b>features</b> |
|------------|-------|--------------|-------|------|-----------------|
|            |       |              |       |      |                 |
|            |       |              |       |      |                 |

- The features column contains, for each document, an array of additional feature scores.
- These can be used for learning and applying LTR models
- Applying an LTR model would override the original score of the document, causing a re-ranking

# Feature Union Operator

## Feature Union (\*\*)

- Say we want to take multiple retrieval approaches as different features. We use  $**$  to denote, that given a set of input documents, use these transformers to obtain additional feature scores

- Formally:  $Rf = (T_1 \ ** \ T_2)(R) :=$

$$R_1 = T_1(R), \quad R_2 = T_2(R)$$

$$Rf = (R_1 \bowtie R_2)[[f_1, f_2] \rightarrow f]$$

- E.g.

BatchRetrieve("BM25") >> ( BatchRetrieve("Tf") \*\* BatchRetrieve("PL2") )

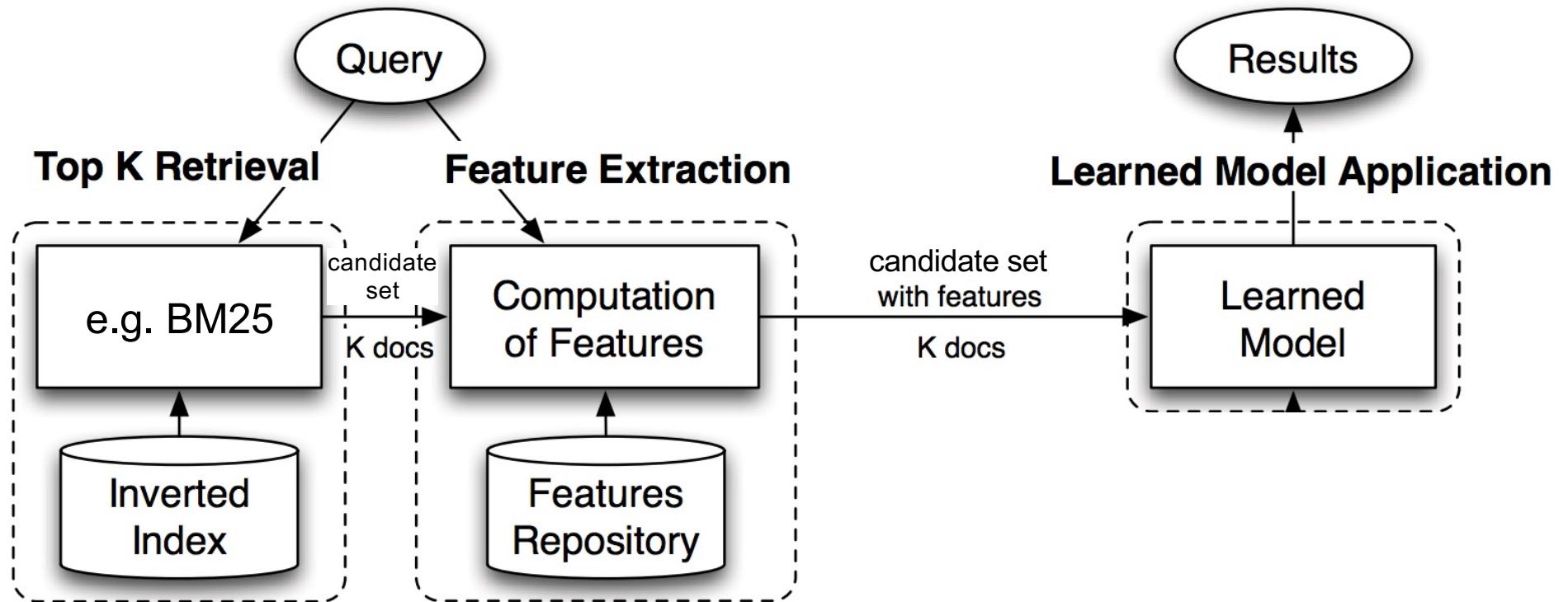
$R$

|  | qid | docno | score              |
|--|-----|-------|--------------------|
|  | 1   | q1    | d5<br>(bm25 score) |

$R_f$

|  | qid | docno | score              | features              |
|--|-----|-------|--------------------|-----------------------|
|  | 1   | q1    | d5<br>(bm25 score) | [tf score, pl2 score] |

# Matching LTR concepts to PyTerrier...

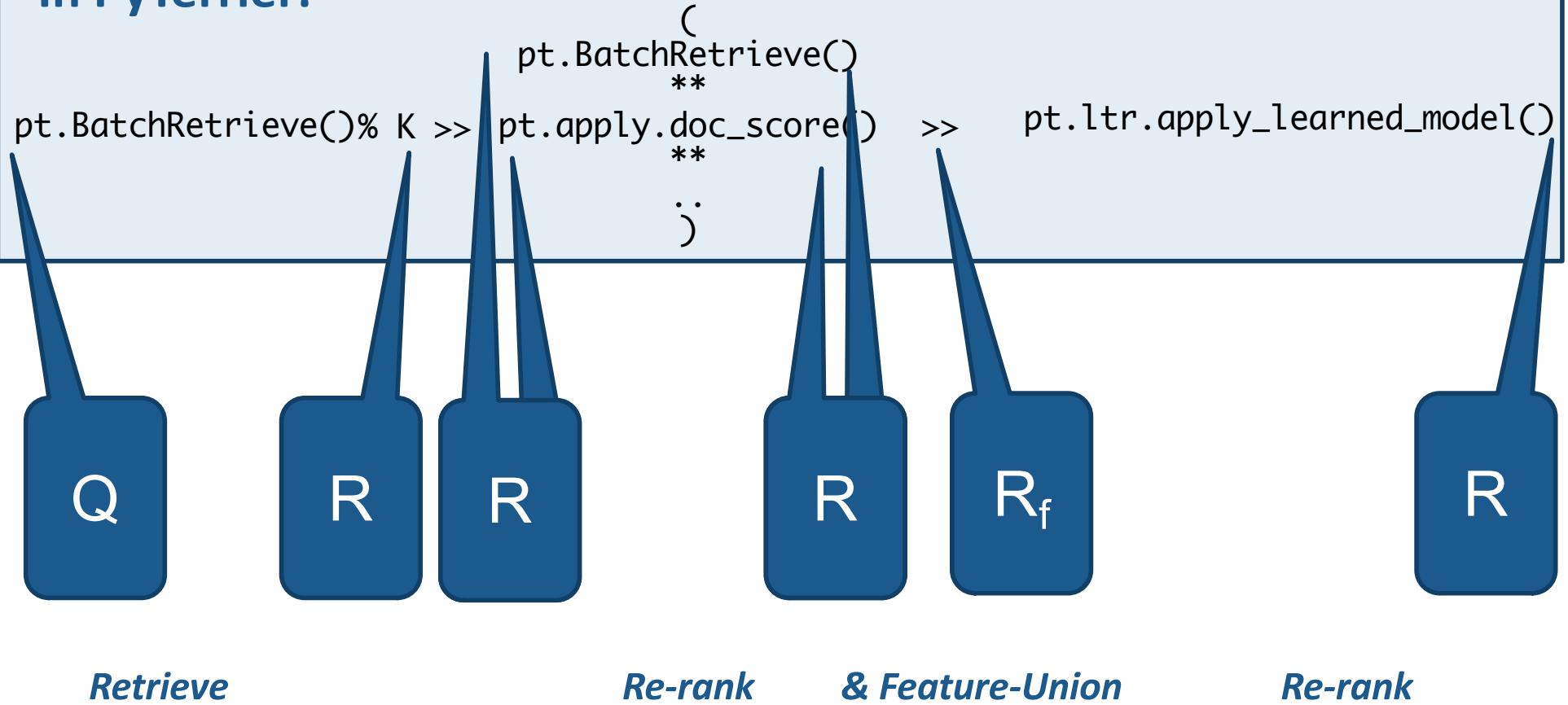


In PyTerrier:

```
pt.BatchRetrieve()  
**  
pt.BatchRetrieve() % K >> pt.apply.doc_score() ** >> pt.ltr.apply_learned_model()  
::
```

# Data Model Transformations

In PyTerrier:



# **Query-Dependent Feature Extraction**



UNIVERSITÀ DI PISA



University  
of Glasgow

BatchRetrieve("BM25") >> ( BatchRetrieve("Tf") \*\* BatchRetrieve("PL2") )

## Here we are using BatchRetrieve as a **re-ranker**

- Feature union is only defined on an input set of documents
- RHS BatchRetrieves will re-score the BM25 retrieved documents using the other weighting models

## We might deploy a number of query dependent features

- Such as additional weighting models, e.g. fields/proximity, that are calculated based on information in the inverted index
- Each BatchRetrieve causes another access to the inverted index posting lists

## FeatureBatchRetrieve is an alternative for Terrier

```
pt.FeatureBatchRetrieve(index, wmodel="BM25",
                        features=[“WMODEL:Tf”, “WMODEL:PL2”])
```

- Multiple features scored in a single pass of the inverted index
- PyTerrier can perform this optimisation when .compile() is called



University  
of Glasgow

UNIVERSITÀ DI PISA

CIKM  
2021  
1-5 NOVEMBER

# TYPES OF MODELS FOR LTR

# **Types of Learned Models (1)**

## **Linear Model**



UNIVERSITÀ DI PISA



### **Linear Models (the most intuitive to comprehend)**

- Many learning to rank techniques generate a linear combination of feature values:

$$score(d, Q) = \sum_f w_f \cdot value_f(d)$$

- Linear Models make some assumptions:
  - **Feature Usage**: They assume that the same features are needed by all queries
  - **Model Form**: The model is only a linear combination of feature values.
    - Contrast this with genetic algorithms, which can learn functional combinations of features, by randomly introducing operators (e.g. try divide feature  $a$  by feature  $b$ ), but are unpractical to learn

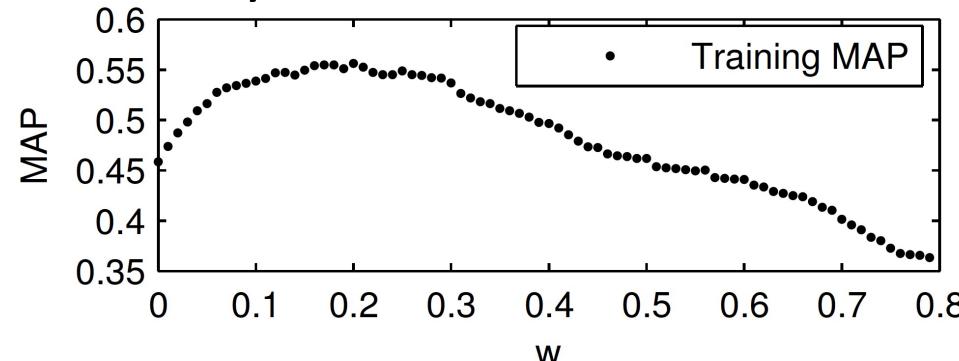
**It is difficult to find  $w_f$  values that maximise the performance of an IR evaluation metric, as they are non-smooth and non-differentiable**

- Typically, techniques such as simulated annealing or stochastic gradient descent are used to empirically obtain  $w_f$  values

# Difficulty in Training

IR evaluation measures are not continuous, and hence are not differentiable wrt. a feature weight

- i.e. they are not smooth as we vary a feature weight  $w$



Why?

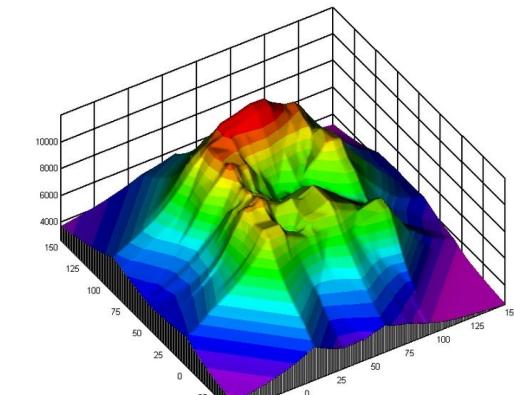


Figure 2: Tuning  $w$ , for  $\log$  PageRank combination.

Metrics don't respond to changes in the scores of documents, but only when a swap between the ordering of two documents takes place

- We need advanced gradient descent/hill-climbing/simulated annealing
- Many of the advances in learning to rank try to address this

Figures from: (a) Craswell et al., SIGIR 2005 (b) Hongning Wang

# (Greedy) Automatic Feature Selection (AFS)

Consider a linear model:

$$score(d, Q) = \sum_f w_f \cdot value_f(d)$$

AFS: Select a  $w_f > 0$  for the feature  $f$  that best improves an evaluation measure  $M$

|       | $f_1$ | $f_2$ | $f_3$ | $f_4$ |
|-------|-------|-------|-------|-------|
| $d_1$ | 0.2   | 0.1   | 0.5   | 0.8   |
| $d_2$ | 0.5   | 0.7   | 0.0   | 0.2   |
| $d_3$ | 0.3   | 0.2   | 0.5   | 0.0   |

Step 1: Select the most effective single feature according to  $M$

|    |      |
|----|------|
| f1 | 0.30 |
| f2 | 0.35 |
| f3 | 0.05 |
| f4 | 0.10 |

→  $(f2, 1)$

| $w_1$ | $w_2$ | $w_3$ | $w_4$ |
|-------|-------|-------|-------|
|       |       |       |       |

Current Best  $M = 0$

# (Greedy) Automatic Feature Selection (AFS)

Consider a linear model:

$$score(d, Q) = \sum_f w_f \cdot value_f(d)$$

AFS: Select a  $w_f > 0$  for the feature  $f$  that best improves an evaluation measure  $M$

|       | $f_1$ | $f_2$ | $f_3$ | $f_4$ |
|-------|-------|-------|-------|-------|
| $d_1$ | 0.2   | 0.1   | 0.5   | 0.8   |
| $d_2$ | 0.5   | 0.7   | 0.0   | 0.2   |
| $d_3$ | 0.3   | 0.2   | 0.5   | 0.0   |

Step 2: Select the feature and corresponding  $w_f$  that most improves  $M$  (e.g. using sim. annealing)

|             |  |
|-------------|--|
| f2 + ? x f1 |  |
| f2 + ? x f3 |  |
| f2 + ? x f4 |  |

| $w_1$ | $w_2$ | $w_3$ | $w_4$ |
|-------|-------|-------|-------|
|       | 1     |       |       |

Current Best  $M = 0.35$

# (Greedy) Automatic Feature Selection (AFS)

Consider a linear model:

$$score(d, Q) = \sum_f w_f \cdot value_f(d)$$

AFS: Select a  $w_f > 0$  for the feature  $f$  that best improves an evaluation measure  $M$

|       | $f_1$ | $f_2$ | $f_3$ | $f_4$ |
|-------|-------|-------|-------|-------|
| $d_1$ | 0.2   | 0.1   | 0.5   | 0.8   |
| $d_2$ | 0.5   | 0.7   | 0.0   | 0.2   |
| $d_3$ | 0.3   | 0.2   | 0.5   | 0.0   |

Step 2: Select the feature and corresponding  $w_f$  that most improves  $M$  (e.g. using sim. annealing)

|                        |      |
|------------------------|------|
| $f_2 + 1.3 \times f_1$ | 0.36 |
| $f_2 + 0.3 \times f_3$ | 0.40 |
| $f_2 + 0.9 \times f_4$ | 0.34 |

→  $(f_3, 0.3)$

| $w_1$ | $w_2$ | $w_3$ | $w_4$ |
|-------|-------|-------|-------|
|       | 1     | 0.3   |       |

Current Best  $M = 0.40$

# (Greedy) Automatic Feature Selection (AFS)

Consider a linear model:

$$score(d, Q) = \sum_f w_f \cdot value_f(d)$$

AFS: Select a  $w_f > 0$  for the feature  $f$  that best improves an evaluation measure  $M$

|       | $f_1$ | $f_2$ | $f_3$ | $f_4$ |
|-------|-------|-------|-------|-------|
| $d_1$ | 0.2   | 0.1   | 0.5   | 0.8   |
| $d_2$ | 0.5   | 0.7   | 0.0   | 0.2   |
| $d_3$ | 0.3   | 0.2   | 0.5   | 0.0   |

Step 2: Select the feature and corresponding  $w_f$  that most improves  $M$  (e.g. using sim. annealing)

|                        |      |
|------------------------|------|
| $f_2 + 1.3 \times f_1$ | 0.36 |
| $f_2 + 0.3 \times f_3$ | 0.40 |
| $f_2 + 0.9 \times f_4$ | 0.34 |

→  $(f_3, 0.3)$

Step 3: Goto step 2, until we observe no more improvements in  $M$ , or no more features

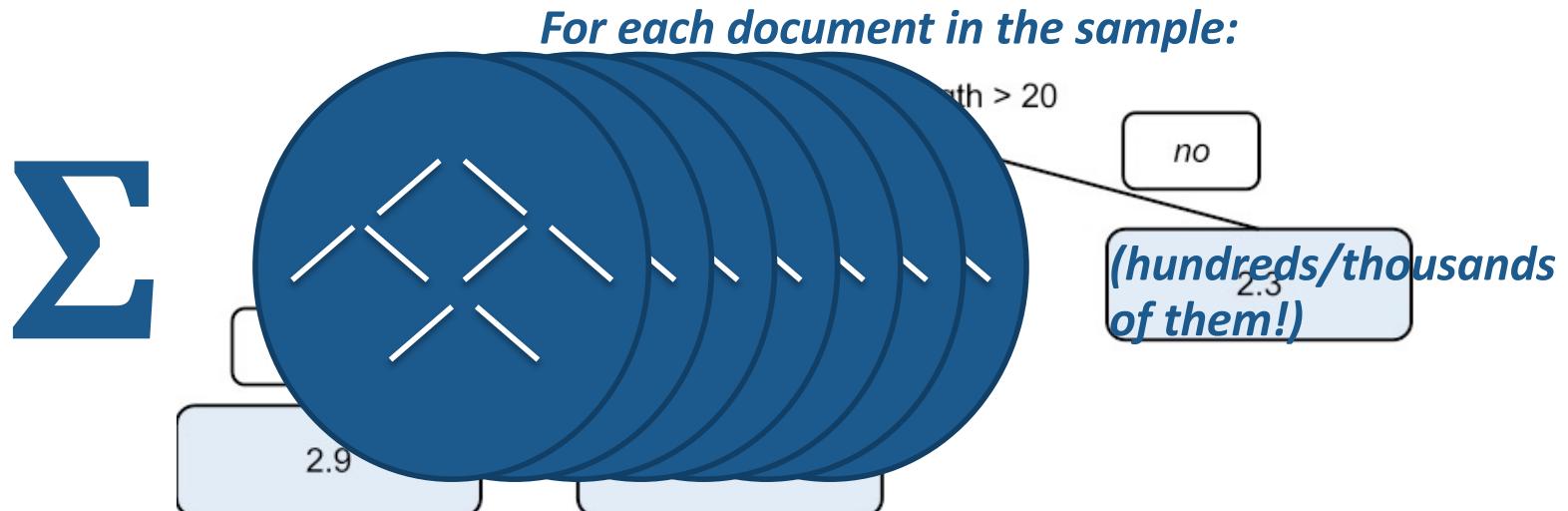
| $w_1$ | $w_2$ | $w_3$ | $w_4$ |
|-------|-------|-------|-------|
|       | 1     | 0.3   |       |

Current Best  $M = 0.40$

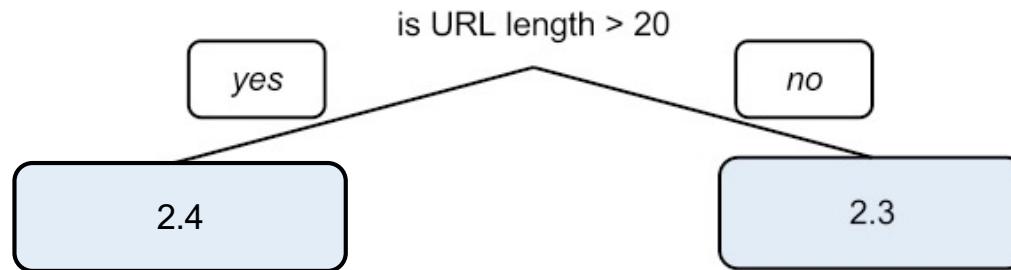
# Type of Learned Models (2)

## Regression Trees

- A *regression tree* is series of decisions, leading to a partial score output
- The outcome of the learner is a “forest” of many such trees, used to calculate the final score of a document for a query
- Their ability to customise branches makes them more effective than linear models
- Regression trees are **pointwise** in nature, but several major search engines have created adapted regression tree techniques that are **listwise**
  - E.g. Microsoft’s **LambdaMART** is at the heart of the Bing search engine



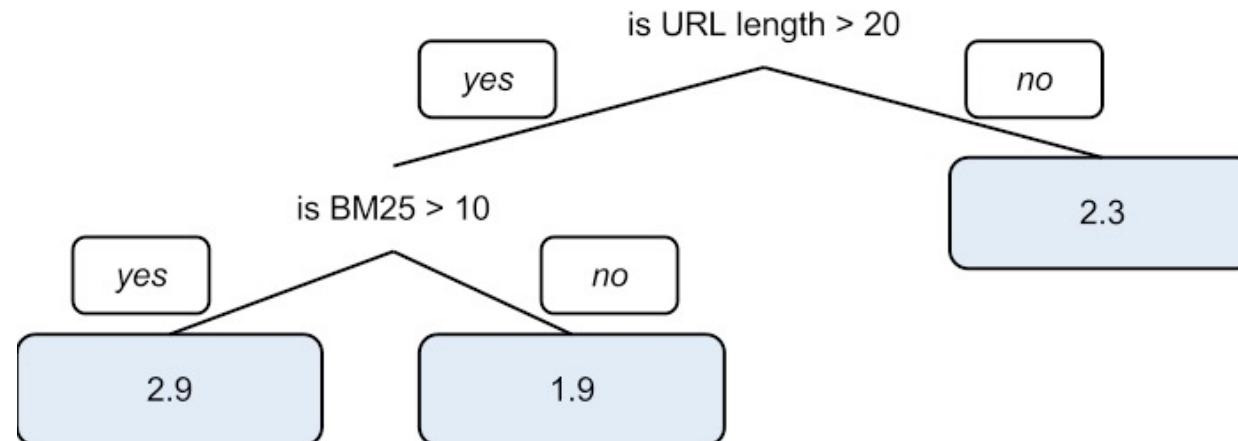
# Growing a Tree



|       | $f_1$ | $f_2$ | $f_3$ | $f_4$ | $L$ |
|-------|-------|-------|-------|-------|-----|
| $d_1$ | 0.2   | 0.1   | 0.5   | 0.8   | 1   |
| $d_2$ | 0.5   | 0.7   | 0.0   | 0.2   | 2   |
| $d_3$ | 0.3   | 0.2   | 0.5   | 0.0   | 0   |

$$\text{Loss} = 1.4$$

*Q: Can we add another decision point?*



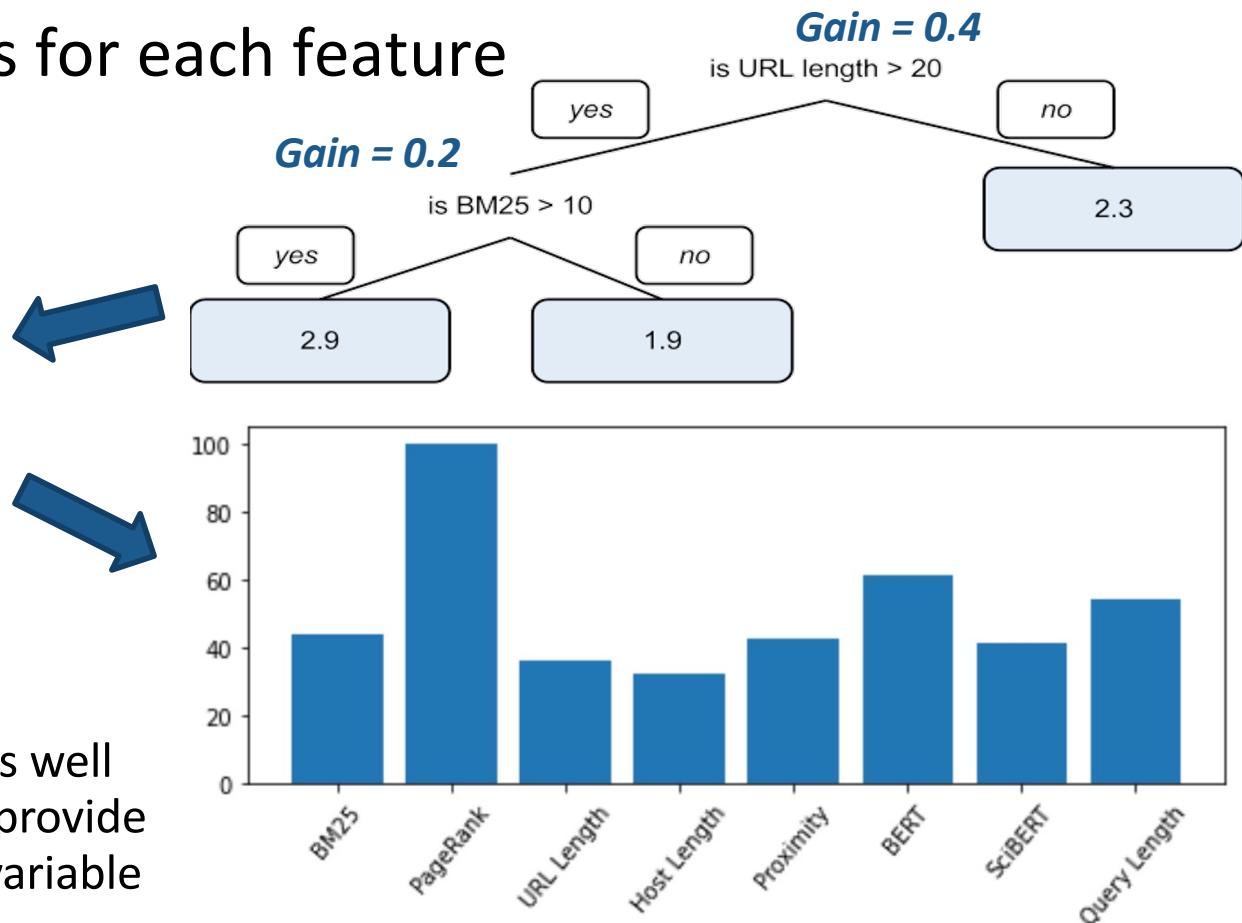
$$\text{Loss} = 1.2 \quad \Rightarrow \text{Gain} = 0.2$$

# Feature Importance

## Regression Trees have an in-built measure of feature importance

- By recording the “gain” at each node during training time
- Summing the gains for each feature

| Feature    | Norm. Total Gain |
|------------|------------------|
| BM25       | 45               |
| PageRank   | 100              |
| URL Length | 39               |
| ...        | ...              |



NB: sklearn’s RandomForest, as well as XGBoost and LightGBM, all provide a `feature_importances_` variable

# **Learning and Applying a Learned Model**



UNIVERSITÀ DI PISA



University  
of Glasgow

## **PyTerrier has transformers that accept standard learners**

- Accessed via `pt.ltr.apply_learned_model()`
- E.g. sci-kit learn has a random forest (pointwise) learner

```
from sklearn.ensemble import RandomForestRegressor
rf = RandomForestRegressor(n_estimators=400)
rf_pipe = pipeline >> pt.ltr.apply_learned_model(rf)
rf_pipe.fit(train_topics, qrels)
```

- Also for XGBoost and LightGBM, which have LambdaMART (listwise) implementations

```
import xgboost as xgb
Lmart = xgb.sklearn.XGBRanker(objective='rank:ndcg')
lmart_pipe = pipeline >> pt.ltr.apply_learned_model(lmart_x, form="ltr")
lmart_pipe.fit(train_topics, train_qrels,
               validation_topics, validation_qrels)
```

- And for FastRank, which has coordinate ascent (linear, listwise) and random forest (pointwise) learners

<https://pyterrier.readthedocs.io/en/latest/ltr.html>



# The Importance of Candidate Set Size

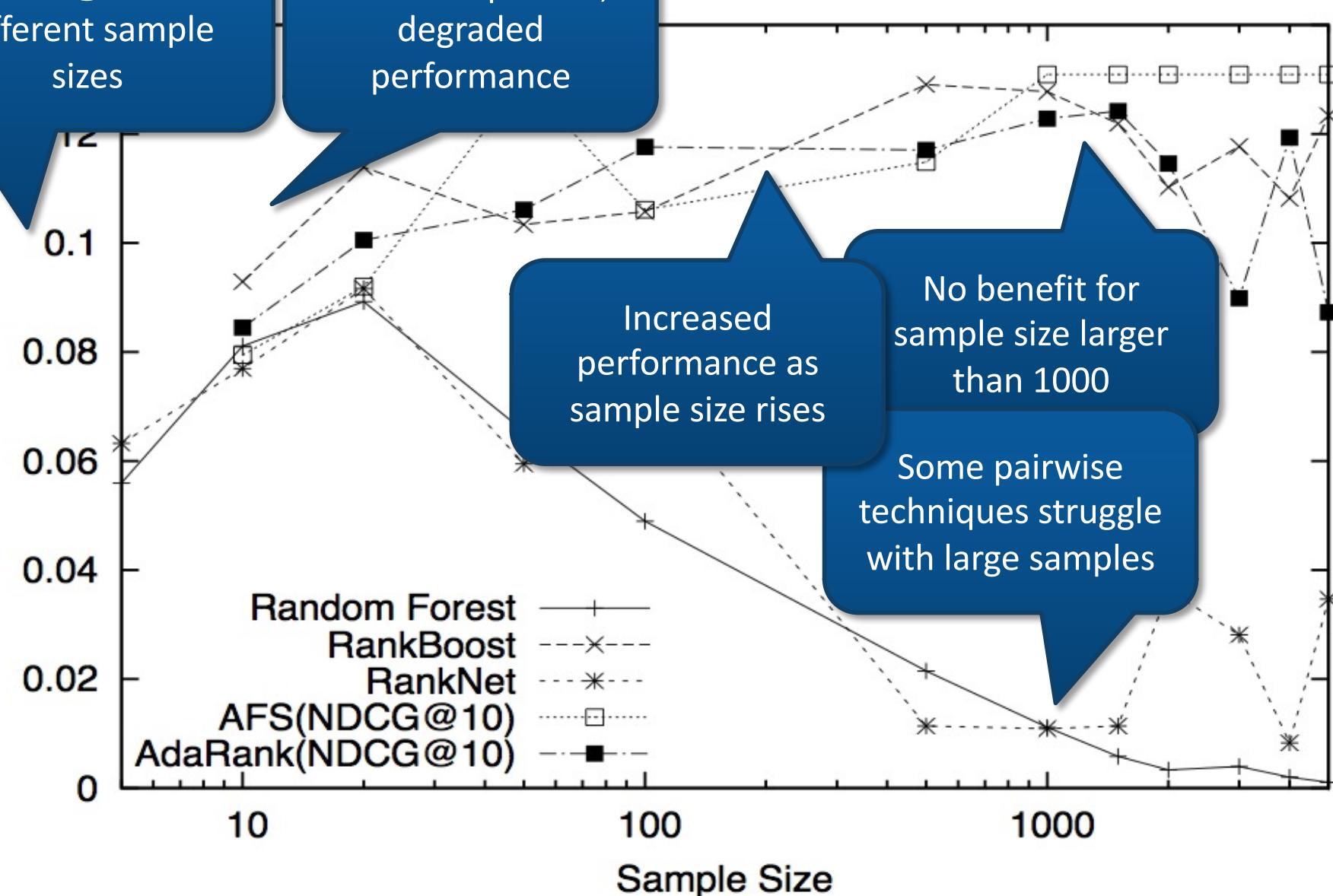
NDCG@20 for different sample sizes

Small sample size, degraded performance

Increased performance as sample size rises

No benefit for sample size larger than 1000

Some pairwise techniques struggle with large samples



# Analysing a Model

Many tree-based learners provide a `feature_importances_` variable

- This allows the researcher to see the importance of various features during *training*
- (How much each feature was used to reduce the loss during tree construction)

Alternatively, utility on the *test* set can be measured by ranking by each feature alone, or ablating (removing) a feature before learning

- E.g. `pt.ltr.ablate_features(int)` is a transformer that zeros out one or more features.
- This is useful when your pipeline is already defined.

```
# learn a model for 3 features, removing one each time
for fid in range(numf):
    ablated = pipeline >> pt.ltr.ablate_features(fid) >> pt.ltr.apply_learned_model(RandomForestRegressor(n_estimators=400))
    ablated.fit(trainTopics, trainQrels, validTopics, validQrels)
    rankers.append(full)
```

- You will see another analysis in the Part 2 notebook

# **PyTerrier Learning to Rank Summary**



UNIVERSITÀ DI PISA



University  
of Glasgow

**Key Operators:** `>>`, `**`

**Key Transformers:**

- Retrieval: `pt.BatchRetrieve`
- Features: `pt.apply.doc_score()`, `pt.FeaturesBatchRetrieve`, `pt.BatchRetrieve`
- Learning to Rank: `pt.ltr.apply_learned_model()`
- Analyse: `pt.ltr.ablate_features()`, `pt.ltr.feature_to_score()`

**LTR Integrations:**

- sklearn: Regression (esp. Random Forests)
- LightGBM: Regression trees/LambdaMART
- xgBoost: Regression trees/LambdaMART
- FastRank: Coordinate Ascent (aka AFS, linear), Random Forests



<https://pyterrier.readthedocs.io/en/latest/ltr.html>

## **LTR Summary**



**Learning to rank is still an important method of integrating many features within a machine-learned search architecture**

- Neural re-ranking models can also fit in this architecture
- Indeed, we found them to be the most important feature for LTR models learned on MSMARCO

**PyTerrier aims to make it as easy as possible to express new features, and analyse the generated models**

- We include integrations of various LTR libraries
- Tensorflow Ranking is one notable possible future integration, which blurs the lines between LTR and neural re-ranking

# References

- [Burges2010] Christopher J.C. Burges. 2010. From RankNet to LambdaRank to LambdaMART: An Overview. Technical Report MSR-TR-2010-82.
- [Chen2016] Tianqi Chen and Carlos Guestrin. 2016. XGBoost: A scalable tree boosting system. In Proceedings of SIGKDD.
- [Foley2019] FastRank alpha, John Foley <https://jjfoley.me/2019/10/11/fastrank-alpha.html>
- [Ke2017] Guolin Ke, Qi Meng, Thomas Finley, Taifeng Wang, Wei Chen, Weidong Ma, Qiwei Ye, and Tie-Yan Liu. 2017. LightGBM: A highly efficient gradient boosting decision tree. Proceedings of NeurIPS.
- [Liu2009] Tie-Yan Liu (2009), "Learning to Rank for Information Retrieval", Foundations and Trends® in Information Retrieval: Vol. 3: No. 3, pp 225-331.
- [Macdonald2012] The Whens and Hows of Learning to Rank. Craig Macdonald, Rodrygo Santos and Iadh Ounis. Information Retrieval 16(5):584-628.
- [Macdonald2012a] On the Usefulness of Query Features for Learning to Rank. Craig Macdonald, Rodrygo Santos and Iadh Ounis. In Proceedings of CIKM 2012.
- [Macdonald2013] About Learning Models with Multiple Query Dependent Features. Craig Macdonald, Rodrygo L.T. Santos, Iadh Ounis and Ben He. *Transactions on Information Systems*. 31(3).
- [Metzler2005] Donald Metzler and W Bruce Croft. 2005. A Markov random field model for term dependencies. In Proceedings of SIGIR.
- [Tonellotto2018] Efficient Query Processing for Scalable Web Search. Nicola Tonellotto, Craig Macdonald and Iadh Ounis. In *Foundations and Trends® in Information Retrieval*. Vol. 12: No. 4-5, pp 319-500, 2018.



University  
of Glasgow

UNIVERSITÀ DI PISA

CIKM  
2021  
1-5 NOVEMBER

Part 2C

## USEFULNESS FOR TEACHING IR

# *Using PyTerrier for Teaching (during a Pandemic)*



UNIVERSITÀ DI PISA



University  
of Glasgow

We (Iadh Ounis, Sean & I) used PyTerrier for teaching IR 2x this year

- All exercises were conducted in Google Colab, and hence suitable for online teaching during the pandemic
- Made use of PyTerrier's Dataset API to provide index/topics/qrels (with authentication for licensed datasets)
- Students prefer notebooks to commandline interfaces (c.f. Terrier)

## Pedagogical appeal

- Full pipeline easily available – indexing, retrieval etc.
- Concepts taught in the course can be easily connected to Python (transformer) classes w/ clear inputs & outputs
  - e.g., a pseudo-relevance feedback component operates after an initial retrieval, and before another ranking component is deployed (final ranking of documents).

# *Practical Courseworks (1/2)*

**Our practical assessments balance ‘programming independence’ with easily markable assessments that scale to large classes**

## **Coursework 1:**

- Perform indexing & retrieval; PyTerrier operators
- Conduct experiments on several query sets, using several classical and probabilistic IR weighting models, e.g., TF-IDF, BM25, PL2, with and without PRF, including applying significance testing, drawing interpolated precision-recall graphs & per-query performance analysis
- Implement a word2vec based QE technique and compare to classical QE

`pt.DFIndexer()`  
`pt.BatchRetrieve()`

`pt.BatchRetrieve()`  
`Bo1QueryExpansion()`  
`pt.Experiment()`

`pt.apply.query()`

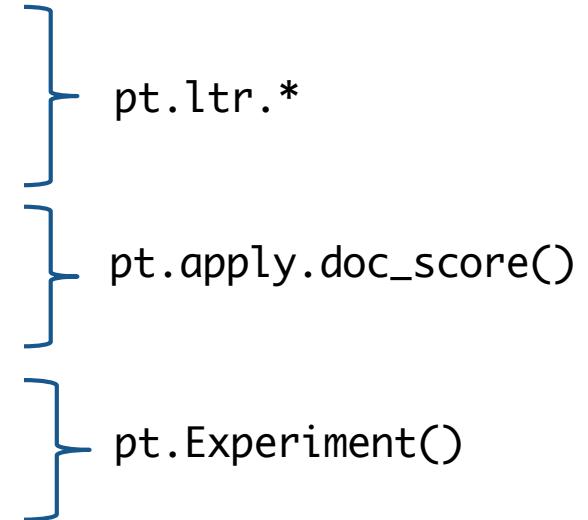
**Most of the exercise has known correct answers can be easily automatically marked in a quiz**

# Practical Courseworks (2/2)



## Coursework 2:

- Train a learning to rank model for web search
- Analyse feature importance
- Define and implement features for URL length and query term proximity
- Conduct experiments, perform significance testing & failure analysis



## Overall feedback:

- PyTerrier builds on familiar technologies: Python, Pandas
- Easy to design and implement new techniques / features
- Students who embrace declarative thinking will write elegant, clear code
- Providing Colab template notebooks helped students to get going
- Increased student performance viz. comparative exercises on commandline/Java

# *Teaching Users*



Delft University of Technology



THE UNIVERSITY  
OF QUEENSLAND  
AUSTRALIA



University  
of Glasgow



McGill  
UNIVERSITY



University  
of Glasgow

UNIVERSITÀ DI PISA

CIKM  
2021  
1-5 NOVEMBER

**Part 2D**  
**WRAPUP**

# Achievements in Part 2



We have **reviewed modern IR architectures**, namely...

- ...how existing IR techniques fit into the PyTerrier datamodel and transformers
- ...how transformer pipelines can be created using operators
- the learning-to-rank architecture, as well as common learners, and how to easily create features using LTR

Finally, we also discussed the usefulness of PyTerrier for teaching  
In the practical session, you will have a chance to **experience** these  
techniques in practice on TREC Covid test collection.

In Part 3, we will **review** contemporary IR re-ranking architectures based on neural networks, as well allowing you to **experience** these through PyTerrier



University  
of Glasgow



CIKM  
2021  
1-5 NOVEMBER

# QUESTIONS?

# A Tutorial Scheduled over 22 hours



University  
of Glasgow

## Run 1 – Live Lectures, Live Labs

Run 1: Aligned with Europe, Africa, and West/South/Central Asia, partially accessible to East Asia/Oceania

| London (GMT) | Content       | Live | Johannesburg (GMT+2) | Delhi (GMT+5:30) | Beijing (GMT+8) | Brisbane (GMT+10) |
|--------------|---------------|------|----------------------|------------------|-----------------|-------------------|
| 0900-1000    | part 1 slides | ✓    | 1100-1200            | 1430-1530        | 1700-1800       | 1900-2000         |
| 1000-1030    | part 1 lab    | ✓    | 1200-1230            | 1530-1600        | 1800-1830       | 2000-2030         |
| 1030-1100    | break         | ☕    | 1230-1300            | 1600-1630        | 1830-1900       | 2030-2100         |
| 1100-1200    | part 2 slides | ✓    | 1300-1400            | 1630-1730        | 1900-2000       | 2100-2200         |
| 1200-1230    | part 2 lab    | ✓    | 1400-1430            | 1730-1800        | 2000-2030       | 2200-2230         |
| 1230-1330    | break         | ☕    | 1430-1530            | 1800-1900        |                 |                   |
| 1330-1430    | part 3 slides | ✓    | 1530-1630            | 1900-2000        |                 |                   |
| 1430-1500    | part 3 lab    | ✓    | 1630-1700            | 2000-2030        |                 |                   |
| 1500-1530    | break         | ☕    | 1700-1730            | 2030-2100        |                 |                   |
| 1530-1630    | part 4 slides | ✓    | 1730-1830            | 2100-2200        |                 |                   |
| 1630-1700    | part 4 lab    | ✓    | 1830-1900            | 2200-2230        |                 |                   |

## Run 2 – Recorded Lectures, Live Labs

Run 2: Aligned with North/South America, partially accessible to East Asia/Oceania

| LA (GMT-7) | Content       | Live | New York (GMT-4) | São Paulo (GMT-3) | Beijing (GMT+8)   | Brisbane (GMT+10) |
|------------|---------------|------|------------------|-------------------|-------------------|-------------------|
| 1100-1200  | part 1 slides | 🎥    | 1400-1500        | 1500-1600         |                   |                   |
| 1200-1230  | part 1 lab    | ✓    | 1500-1530        | 1600-1630         |                   |                   |
| 1230-1300  | break         | ☕    | 1530-1600        | 1630-1700         |                   |                   |
| 1300-1400  | part 2 slides | 🎥    | 1600-1700        | 1700-1800         |                   |                   |
| 1400-1430  | part 2 lab    | ✓    | 1700-1730        | 1800-1830         |                   |                   |
| 1430-1530  | break         | ☕    | 1730-1830        | 1830-1930         |                   |                   |
| 1530-1630  | part 3 slides | 🎥    | 1830-1930        | 1930-2030         | 0630-0730 (6 Nov) | 0830-0930 (6 Nov) |
| 1630-1700  | part 3 lab    | ✓    | 1930-2000        | 2030-2100         | 0730-0800 (6 Nov) | 0930-1000 (6 Nov) |
| 1700-1730  | break         | ☕    | 2000-2030        | 2100-2130         | 0800-0830 (6 Nov) | 1000-1030 (6 Nov) |
| 1730-1830  | part 4 slides | 🎥    | 2030-2130        | 2130-2230         | 0830-0930 (6 Nov) | 1030-1030 (6 Nov) |
| 1830-1900  | part 4 lab    | ✓    | 2130-2200        | 2230-2300         | 0930-1000 (6 Nov) | 1030-1200 (6 Nov) |

Breakout Lab Facilitators: Sean, Nic, Craig

Breakout Lab Facilitators: Eugene, Luca

# *Practical Time*



The tutorial Github repo has links to the notebook for  
**Part 2**

- <https://github.com/terrier-org/cikm2021tutorial>
- Press the  Open in Colab link for each notebook to start a Colab session

**Timings:**

- Practical – in breakout rooms – until 1230
- Lunch break 1230-1330
- Part 3 resumes at 1330

*Run 1 times, GMT*

If you are leaving us here, please complete our feedback quiz <https://forms.office.com/r/RiYSAxAKhk> [link also on repo]