

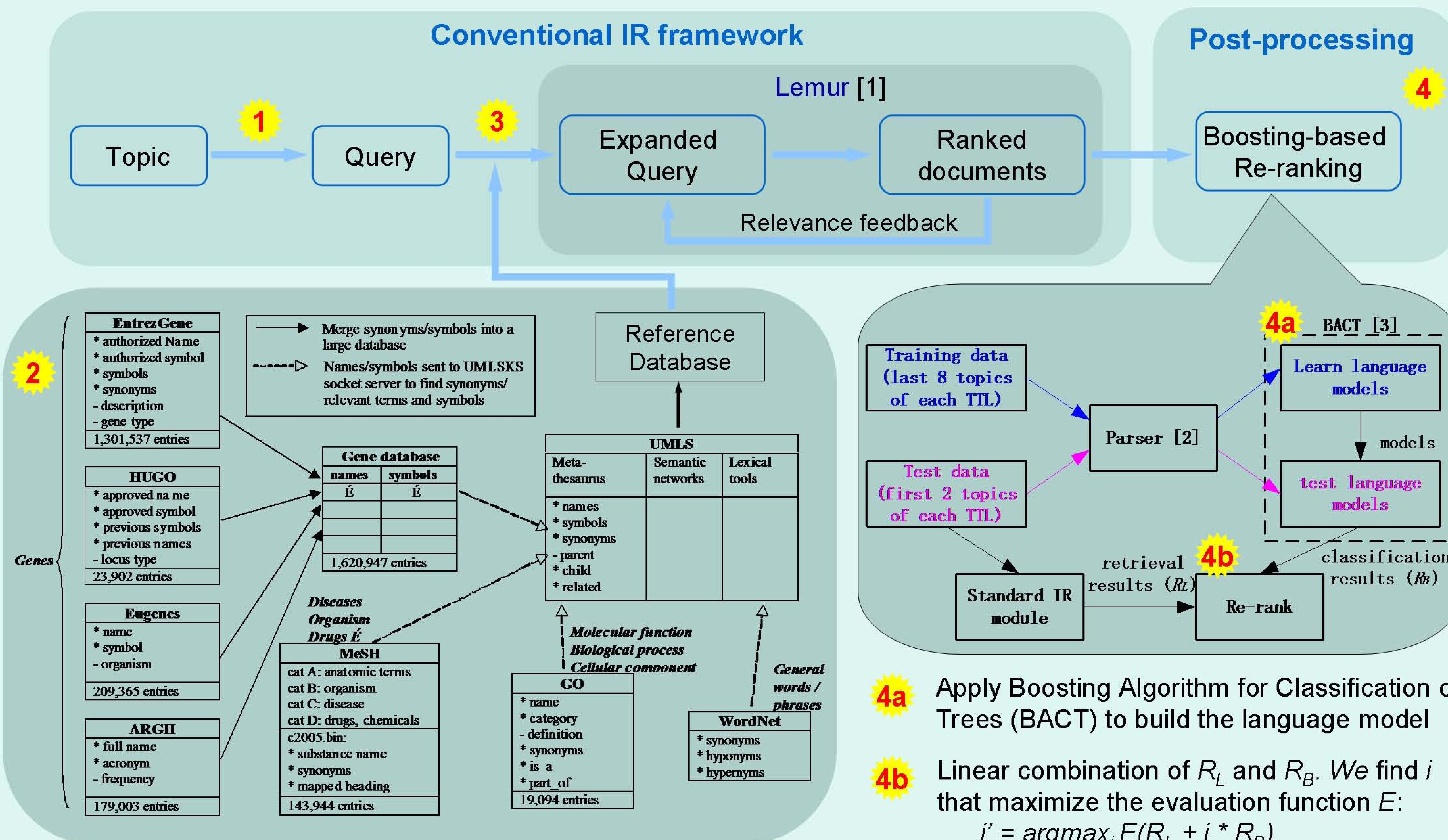
Synonym-based Query Expansion and Boosting-based Re-ranking: A Two-phase Approach for Genomic Information Retrieval

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- 1 Query generation:** manually select keywords from official topic and build the structured query for each topic
- 2 Reference database construction:** collecting synonyms from a large collection of ontology sources: EntrezGene, HUGO, Eugenes, ARGH, MeSH, GO, UMLS and WordNet.
- 3 Query expansion:** look up synonyms of keywords in the reference database
- 4 Re-ranking in the post-processing:** apply a boosting-based classification algorithm to re-rank the retrieved documents

Experimental results:
The boosting-based re-ranking does help when *bpref* of the conventional IR system is low.



| topic # | metrics | $i=0$ (i') | $i=10$ |
|---------|---------|----------------|--------|
| 100 | MAP | 0.2221 | 0.1785 |
| | bpref | 0.8649 | 0.8649 |
| | P10 | 0.4 | 0.3 |
| | P100 | 0.28 | 0.22 |
| 101 | MAP | 0.0685 | 0.0195 |
| | bpref | 0.75 | 0.75 |
| | P10 | 0 | 0 |
| | P100 | 0.07 | 0.07 |

Table 1: Performance of re-ranking on TTL #1

| topic # | metrics | $i=0$ | $i=15$ (i') |
|---------|---------|--------|-----------------|
| 110 | MAP | 0.0012 | 0.0024 |
| | bpref | 0.25 | 0.25 |
| | P10 | 0 | 0 |
| | P100 | 0 | 0.01 |
| 111 | MAP | 0.0492 | 0.1602 |
| | bpref | 0.4356 | 0.4356 |
| | P10 | 0.1 | 0.7 |
| | P100 | 0.1 | 0.4 |

Table 2: Performance of re-ranking on TTL #2

| topic # | metrics | $i=0$ (i') | $i=10$ |
|---------|---------|----------------|--------|
| 120 | MAP | 0.6113 | 0.2410 |
| | bpref | 0.8145 | 0.8145 |
| | P10 | 1 | 0.3 |
| | P100 | 0.88 | 0.29 |
| 121 | MAP | 0.6697 | 0.0328 |
| | bpref | 0.8810 | 0.8810 |
| | P10 | 0.8 | 0 |
| | P100 | 0.34 | 0 |

Table 3: Performance of re-ranking on TTL #3

[1] Lemur. 2005. Language Modeling Toolkit 4.1. <http://www.lemurproject.org>.

[2] Eugene Charniak. A maximum-entropy-inspired parser. NAACL 2000.

[3] Taku Kudo and Yuji Matsumoto. A boosting algorithm for classification of semi-structured text. EMNLP 2004.