

Ranking Biomedical Passages for Relevance and Diversity

University of Wisconsin-Madison at TREC Genomics 2006

Andrew B. Goldberg, David Andrzejewski, Jurgen Van Gael, Burr Settles, Xiaojin Zhu, Mark Craven^{*} Computer Science Department, * Biostatistics Department



Outline

Genomics Track Task Overview
 System Overview
 Reranking using Absorbing Random Walks Results
 Discussion

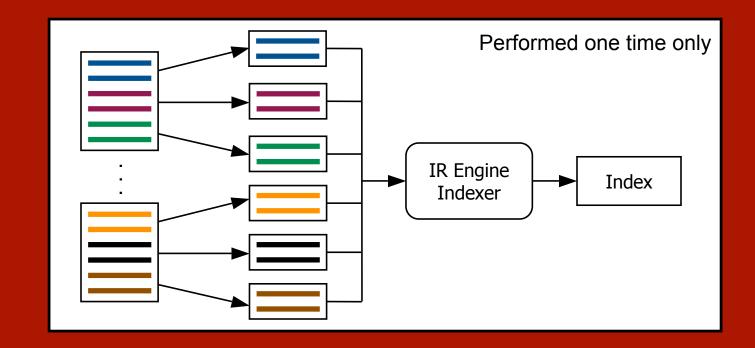
Genomics Track Task Overview

- Given 160,000 full-text biomedical articles
 Given a scientific query:
 - What is the role of PrnP in mad cow disease?
- Find and rank short passages about different aspects of the question
- Document, passage, and aspect-level evaluation metrics
- UW-Madison submitted 3 automatic runs



Phase I: Indexing

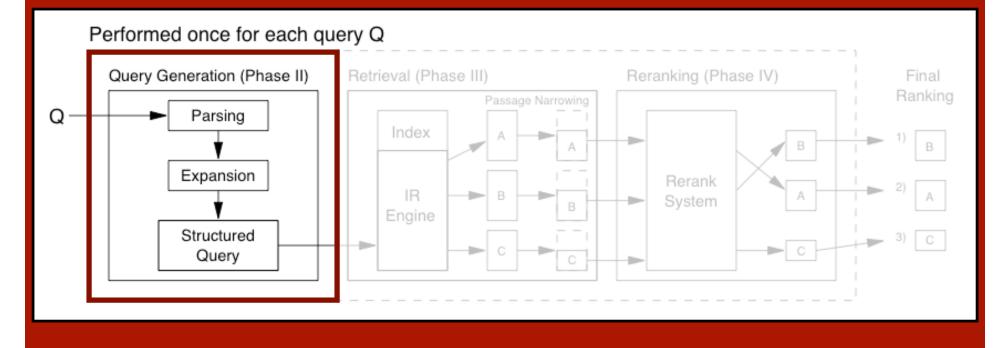
- 1. Split documents into legal spans
- 2. Build index using off-the-shelf IR engine
 - Lemur toolkit with Indri index





Phase II: Query Generation

- 1. Parse natural language topic questions
- 2. Expand queries using online resources
- 3. Automatically generate structured queries





Phase II: Query Generation

Example parse (before stop word removal):

What is the role of PrnP in mad cow disease ?

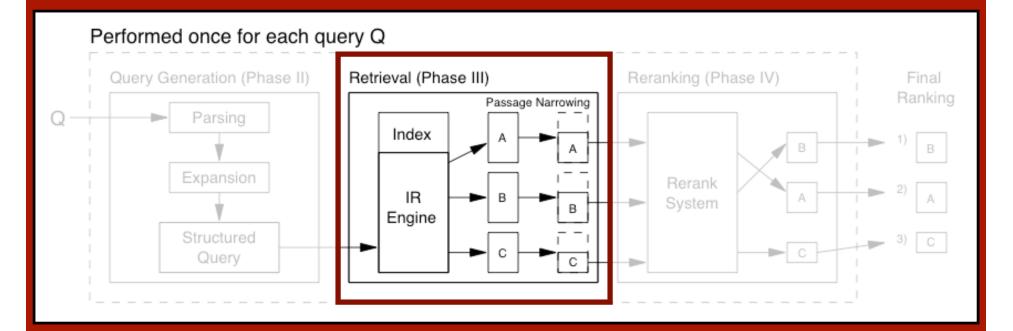
Structured query with expansion terms:

```
#filreq(
    #band(
        #syn( #1( PrnP ) #1( prion protein )...)
        #syn( #1( mad cow disease ) #1( BSE )
            #1( Bovine Spongiform Encephalopathy )...)
)
#combine( #1( PrnP ) #1( prion protein ) ...
#1( mad cow disease ) #1( BSE )
        #1( Bovine Spongiform Encephalopathy )...
)
```



Phase III: Retrieval

 Run query using off-the-shelf IR engine
 Trim paragraphs to create passages that include only the relevant sentences





Phase III: Retrieval

Example of passage narrowing:

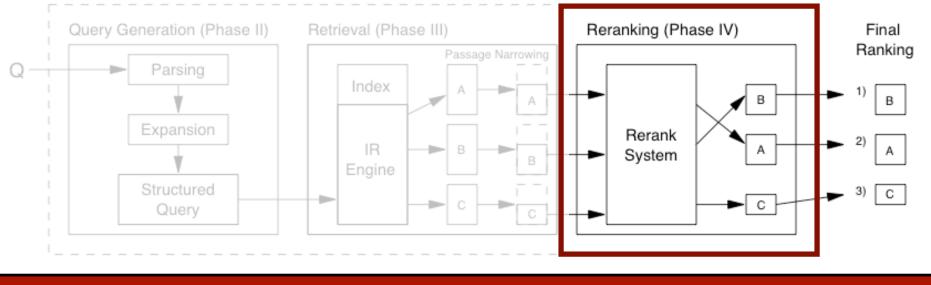
In December 1984 a UK farmer called a veterinary surgeon to look at a cow that was behaving unusually. Seven weeks later the cow died. Early in 1985 more cows from the same herd developed similar clinical signs. In November 1986 **bovine spongiform encephalitis (BSE)** was first identified as a new disease, later reported in the veterinary press as a novel progressive **spongiform encephalopathy**. Later still the causal agent of **BSE** was recognized as an abnormal **prion protein**. Since the outset the story of **BSE** has been beset by problems.



Phase IV: Reranking

Run 1: Indri ranking Run 2: Clustering-based reranking Run 3: Absorbing random walk reranking

Performed once for each query Q





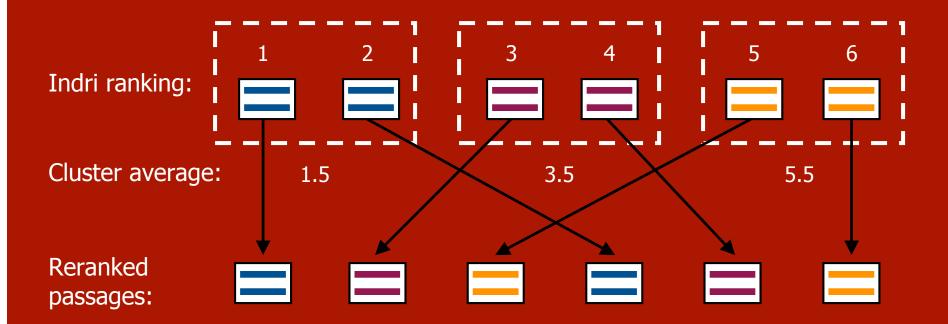
Clustering-Based Reranking

Cluster passages using bag-of-words vectors and cosine similarity
 Assume clusters represent aspects
 Interleave results from clusters to achieve aspect diversity



Clustering-Based Reranking

Example with 3 clusters



(Note: We arbitrarily used 10 clusters in Run 2)

Absorbing Random Walk Reranking

Produces new ranking such that

 A highly ranked passage is *central* to a local group in the set
 Top ranked items cover many *diverse* groups
 Initial ranking is included as prior knowledge

 Achieves these goals using absorbing Markov chain random walks



High-Level View of Algorithm

Random walk on a graph over passages
Ranked passages become absorbing states
Absorbing states "drag down" importance of similar unranked states
Newly ranked states differ from previously ranked states to promote diversity

Algorithm Input

Graph W with n nodes (items to rank) - Represented by *n* x *n* weight matrix Large weight means similar items Prior distribution r based on initial ranking High initial ranks have high probabilities - No prior ranking = uniform distribution (all 1/n) • Weight $\lambda \in [0,1]$ – Balances influence of W versus r



Teleporting random walk
 Random walker moves around graph

 With probability λ:
 Walks to a neighbor state based on edge weights W
 More likely to walk to similar state
 Otherwise:

 Teleports randomly according to r

More likely to walk to state with high initial rank

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Finding the First Item to Rank

1. Create transition matrix by normalizing rows of *W* weight matrix

$$\tilde{P}_{ij} = \frac{w_{ij}}{\sum_{k=1}^{n} w_{ik}}$$

2. Transform into teleporting random walk by interpolating each row with prior *r*

$$P = \lambda \tilde{P} + (1 - \lambda) \mathbf{1} \mathbf{r}^{\mathsf{T}}$$



- Stationary distribution of random walk

 Defines visiting probabilities of nodes

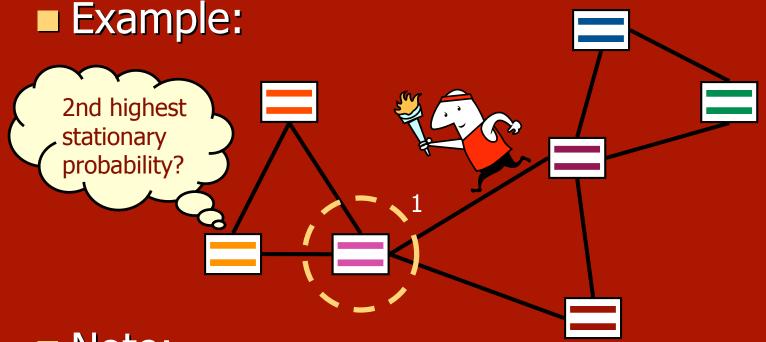
 Dense regions of graph (soft clusters) have high probabilities
 High probability states regarded as central, most important items
 - Like Google's PageRank algorithm



- 3. Find unique stationary distribution $\pi = P^{\top}\pi$
- 4. Select first item as the state with the largest stationary probability

$$g_1 = \operatorname{argmax}_{i=1}^n \pi_i$$





Note:

- Only the larger *W* edge weights are shown
- Interpolation with r makes it fully connected



Centrality versus Diversity

Stationary distribution lacks diversity – High probability items from same local groups To ensure diversity: First ranked item becomes an absorbing state – Walker can fall in "black hole" and walk ends Stationary distribution now uninformative All walks will eventually get absorbed Need alternate way to select items

Ranking the Remaining Items

New selection criterion:

 Expected number of visits before absorption
 How does this promote diversity?

 Walker spends less time in

less time in states with high probability of being absorbed

Walker spends more time in dissimilar regions of graph (before getting absorbed)

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Ranking the Remaining Items

While more items to rank:

- 1. Turn ranked states into absorbing states
- 2. Compute expected number of visits per unranked item
- Select the item with the maximum expected number of visits

Ranking the Remaining Items

Turn ranked states *G* into absorbing states For $g \in G$, $P_{gg} = 1$, $P_{gi} = 0$, $\forall i \neq g$ Arrange *P* with ranked before unranked states:

$$P = \begin{bmatrix} \mathbf{I}_{G} & \mathbf{0} \\ R & Q \end{bmatrix} \leftarrow \text{Ranked items}$$
$$\leftarrow \text{Unranked items}$$

Find fundamental matrix: $N = (\mathbf{I} - Q)^{-1}$

 N_{ij} = expected number of visits to state j before absorption, if the walk started in state i



Ranking the Remaining Items

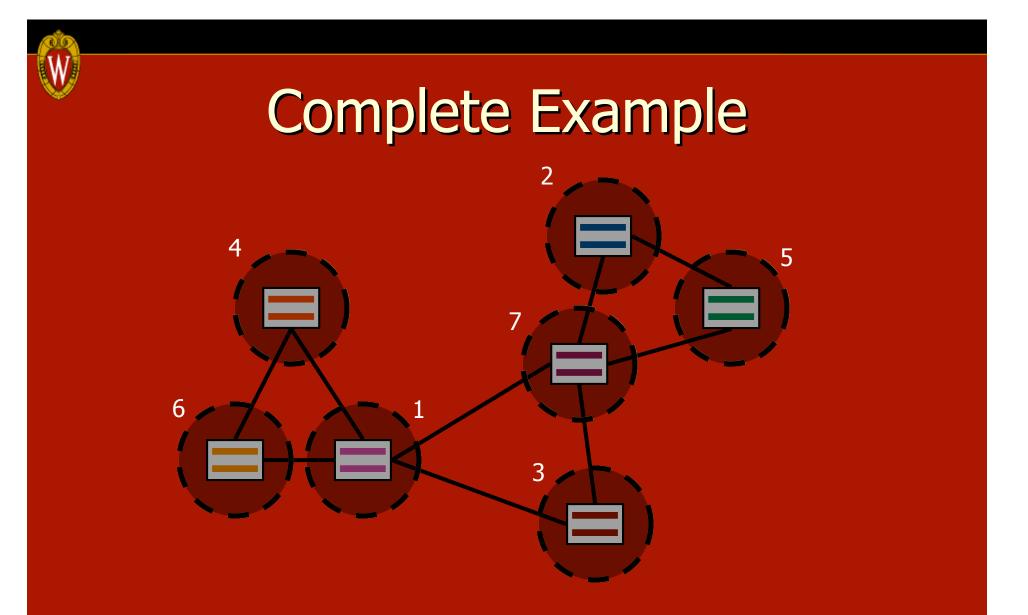
The expected number of visits per state:

 $\mathbf{v} = \frac{N^{\top} \mathbf{1}}{n - |G|}$

 v_i = expected number of visits to state j before absorption, regardless of starting state

Picking the next item:

 $g_{|G|+1} = \arg \max_{i=|G|+1}^{n} v_i$



Notice that the ranking hops between dense regions (hopefully different aspects) in the graph



Parameters in Run 3

Weight matrix W

 Symmetric 10-NN graph using cosine similarity of bag-of-words vectors
 Initial ranking distribution r
 r_i ∝ (n − initialRank_i + 1)

Trade-off parameter λ

 Arbitrarily set to 0.6 to put more emphasis on graph but still have influence of initial ranking



Results

Run	Document	Passage	Aspect
Indri Ranking	0.2368	0.0188	0.1516
Clustering	0.2030	0.0137	0.1319
Random Walk	0.2208	0.0159	0.1411

Mean average precision (MAP) scores

Document and Passage seem mediocre
 Aspect appears competitive, but reranking methods fail to improve over baseline



Discussion

Poor document and passage MAP scores

- Query generation inadequate
- No results produced for some topics
- Perhaps exact matching in queries too strict
- Solutions?
 - Refine parsing technique
 - Use less restrictive query operators
 - Consult additional resources (GO, UMLS, etc)



Reranking Discussion

Irrelevant documents appear diverse

 Incorrectly placed even higher in ranks

 Similarity graph may be inappropriate

 Needs to correlate with aspect similarity
 Could use TF-IDF vectors
 (with IDF based on current set of passages)
 Also, KL-divergence between language models

Thank You!

Questions?