

A Practical Algorithm for Topic Modelling with Provable Guarantees

Sanjeev Arora

Rong Ge

Yoni Halpern

David Mimno

Ankur Moitra

David Sontag

Yihcen Wu

Michael Zhu

Presented by: Vanush Vaswani and Kristy Hughes

1 Introduction

2 Topic Modelling

3 Algorithm

4 Efficiently Finding Anchor Words

5 Topic Recovery via Bayes' Rule

6 Experimental Results

7 Conclusion

Information Overload



Effective Organisation





- ① Introduction
- ② Topic Modelling**
- ③ Algorithm
- ④ Efficiently Finding Anchor Words
- ⑤ Topic Recovery via Bayes' Rule
- ⑥ Experimental Results
- ⑦ Conclusion

Topics

Topics

gene 0.04
dna 0.02
genetic 0.01
...

life 0.02
evolve 0.01
organism 0.01
...

brain 0.04
neuron 0.02
nerve 0.01
...

data 0.02
number 0.02
computer 0.01
...

Topics are distributions over words

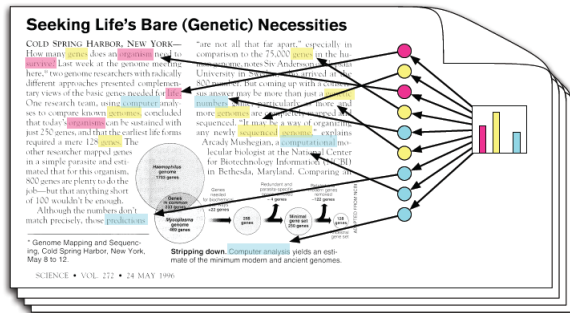


Documents

Documents have
distribution of topics

Documents

Topic proportions and
assignments





Topic Modelling

Topics

gene 0.04
dna 0.02
genetic 0.01
...

life 0.02
evolve 0.01
organism 0.01
...

brain 0.04
neuron 0.02
nerve 0.01
...

data 0.02
number 0.02
computer 0.01
...

Documents

Seeking Life's Bare (Genetic) Necessities

COLD SPRING HARBOR, NEW YORK—How many **genes** does an **organism** need to **survive**? Last week at the genome meeting here,* two genome researchers with radically different approaches presented complementary views of the basic genes needed for **life**. One research team, using **computer** analyses to compare known **genomes**, concluded that today's **organisms** can be sustained with just 250 genes, and that the earliest life forms required a mere 128 **genes**. The other researcher mapped genes in a simple parasite and estimated that for this organism, 800 genes are plenty to do the job—but that anything short of 100 wouldn't be enough.

Although the numbers don't match precisely, those **predictions**

"are not all that far apart," especially in comparison to the 75,000 **genes** in the human genome, notes Siv Anderson, a geneticist at the University in Sydney, who arrived at the 800 number. But coming up with a consensus answer may be more than just a **difficult** numbers game, particularly as more and more **genomes** are completely mapped and sequenced. "It may be a way of organizing any newly **sequenced genome**," explains Arcady Mushegian, a **computational** molecular biologist at the National Center for Biotechnology Information (NCBI) in Bethesda, Maryland. Comparing an



* Genome Mapping and Sequencing, Cold Spring Harbor, New York, May 8 to 12.

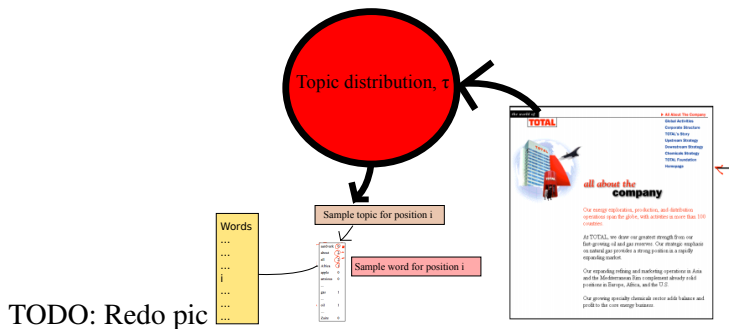
Stripping down. Computer analysis yields an estimate of the minimum modern and ancient genomes.

SCIENCE • VOL. 272 • 24 MAY 1996

Topic proportions and assignments

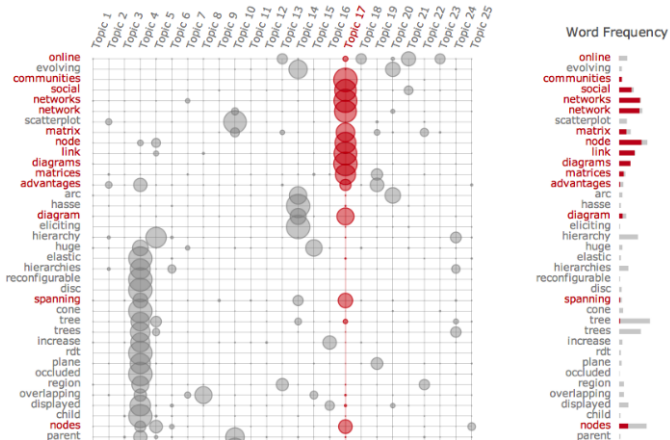
Task

- Assume documents are generated by probabilistic model with unknown variables
- Infer hidden structure onto document
- Situate new document into model



Word-topic Matrix

Extracted: Word-topic matrix



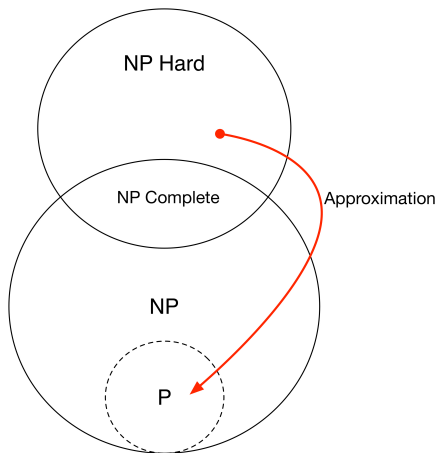
Aim: Find document-topic matrix

Anchor Words

- Word-topic distributions are separable
- There is a word unique to each topic
- Indicates document is partially about that topic
- Can learn parameters in polynomial time provided there is a large enough number of documents

Approximate Inference & Provable Guarantees

- Document-topic inference:
 - NP-hard
- Approximate techniques
- Provably polynomial-time?





- ① Introduction
- ② Topic Modelling
- ③ Algorithm**
- ④ Efficiently Finding Anchor Words
- ⑤ Topic Recovery via Bayes' Rule
- ⑥ Experimental Results
- ⑦ Conclusion

Algorithm

Input: Corpus \mathcal{D} , Number of topics K

Output: Word-topic matrix A , topic-topic matrix R

- 1 Compute word-word co-occurrence matrix
- 2 Normalize the matrix
- 3 Find anchor words
- 4 Recover topics

Assumptions:

- Topics may be correlated
- Word-topic distributions are separable

Contributions

① Anchor Selection

- Combinatorial rather than ILP
- Stable in the presence of noise
- polynomial sample complexity

② Recovery step

- Previous matrix-inversion approach sensitive to noise
- Replaced with Gradient-based inference

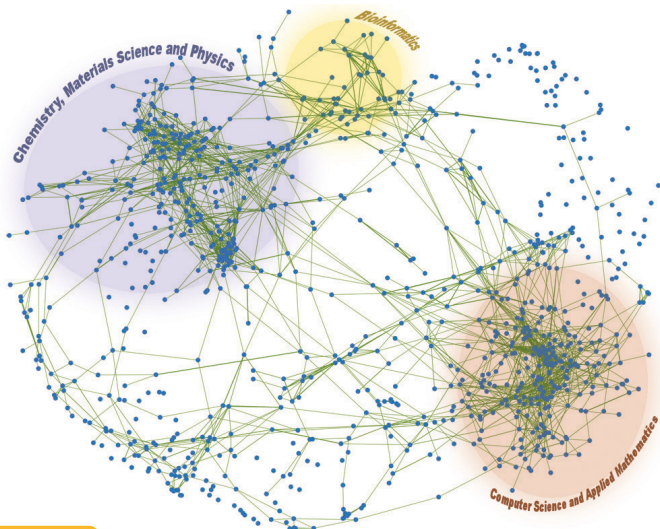
③ Empirical comparison of algorithms

- ① Introduction
- ② Topic Modelling
- ③ Algorithm
- ④ Efficiently Finding Anchor Words**
- ⑤ Topic Recovery via Bayes' Rule
- ⑥ Experimental Results
- ⑦ Conclusion

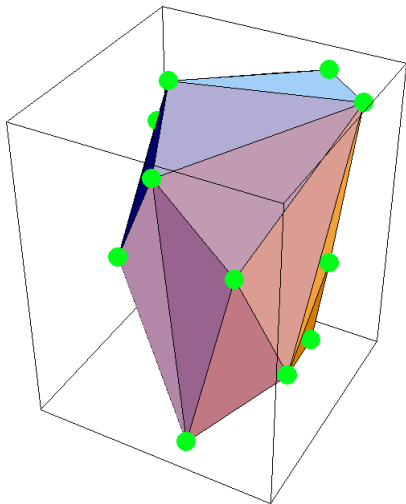
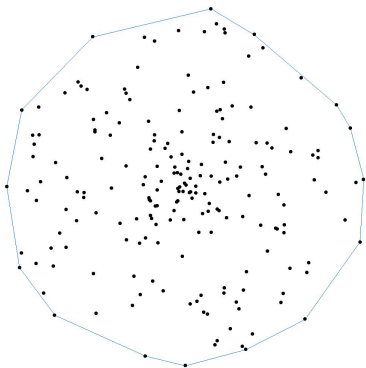
Word-word co-occurrence matrix

	bank	California	Canada	career	careers	employers	employment	federal	human	job	jobs	listings	openings	opportunities	positions	recruiters	resources	resume	resumes	retirement	search	state	texas	unemployment	work
bank	--																								
California	--																								
Canada	1	--																							
career	3	3	--																						
careers			2	9	--																				
employers		2		11	7	--																			
employment	3	26	22	66	10	16	--																		
federal	1	1	5		1		11	--																	
human		4	12	1	1			4	--																
job	34	14	2	49	8	13	92	13	2	--															
jobs		18	6	62	11	27	204	19	2	74	--														
listings		4	2	15	4	9	68	2	55	44		--													
openings		4		7	2	9	28		49	30		--													
opportunities	4	8	3	51	9	13	181	9		84	106	25	19	--											
positions		1		8	2	10	19			16	20	9	13	21	--										
recruiters				10	4	3	9			5	4	2	2	5	2	--									
resources		4	12		1			4	74	3	2						--								
resume		4	3	5		2	3	1	1	10	3			1	2	1		--							
resumes				8	3	3	11			5	16	1		8	5			15	--						
retirement		1	1						3			2		1						--					
search			3	4	6			10			18	6		6	2			3	1		--				
state			4	1			18		1	12	7	6		3			1			2		--			
texas	2			1			18			12	6		1	2						9			--		
unemployment																						2	2	--	
work			2	1		3	3	2		2	8	2	4	7	5				1			1	2	--	

Words as vertices



Convex Hull



Computing Convex Hull

- Efficient for 2 dimensions - $O(n \log n)$
- Inefficient for $n > 2$ dimensions
- Complexity depends on method and approximation used
- Previous method: ILP
- New method: Recursive greedy
 - 1 Compute subspace span of current convex hull
 - 2 Find point furthest from this sub-span
 - 3 Add point to convex hull
 - 4 Repeat until K points found

TODO: Work out how the whole convex hull - words as vertices work. I think what we have here is wrong because there is no approximation



- 1 Introduction
- 2 Topic Modelling
- 3 Algorithm
- 4 Efficiently Finding Anchor Words
- 5 Topic Recovery via Bayes' Rule**
- 6 Experimental Results
- 7 Conclusion

Topic Recovery Task

- Recovers the topics
- Represented as topic-word distributions
- Topic uniquely identified by anchor word

Previous method

- 1 Discard rows not containing anchor words from word-word co-occurrence matrix (Q)
- 2 Permute matrix into a diagonal matrix
- 3 We know that $Q = ARA^T$ where A is the word-topic matrix and R is the topic-topic matrix
- 4 Solve $Q = ARA^T$ using matrix inversion

Word-word co-occurrence probability matrix

For an anchor word (row) in the co-occurrence matrix

$$Q_{s_k,j} = \sum_{k'} p(z_1 = k' | w_1 = s_k) p(w_2 = j | z_1 = k')$$

= 1 because of the anchor word property

$$= p(w_2 = j | z_1 = k) = c_{i,k}$$

For any other row

$$\bar{Q}_{i,j} = \sum_k p(z_1 = k | w_1 = i) p(w_2 = j | z_1 = k)$$

But this is clearly a convex combination of anchor words

$$\bar{Q}_{i,j} = \sum_k c_{i,k} \bar{Q}_{s,k}$$



New method

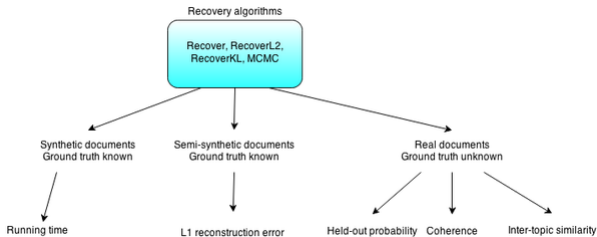
- Row normalize Q into \bar{Q}
- Recover A and R using Bayes rule

$$p(w_1 = i | z_1 = k) = \frac{p(z_1 = k | w_1 = i)p(w_1 = i)}{\sum_i p(z_1 = k | w_1 = i')p(w_1 = i')}$$



- ① Introduction
- ② Topic Modelling
- ③ Algorithm
- ④ Efficiently Finding Anchor Words
- ⑤ Topic Recovery via Bayes' Rule
- ⑥ Experimental Results**
- ⑦ Conclusion

Experiments



Metrics

- Ground truth known
 - Reconstruction error
- Ground truth unknown
 - Held out probability - probability of an unseen document under the learned model
 - Coherence - measure of semantic quality
 - Inter-topic similarity - measure of uniqueness of topics

Results - Efficiency

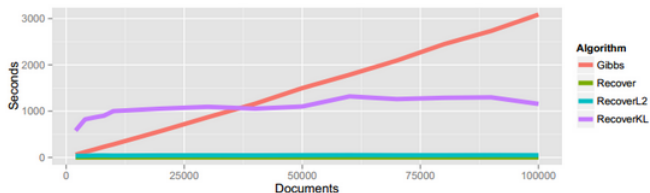


Figure 1. Training time on synthetic NIPS documents.

Results - Reconstruction error

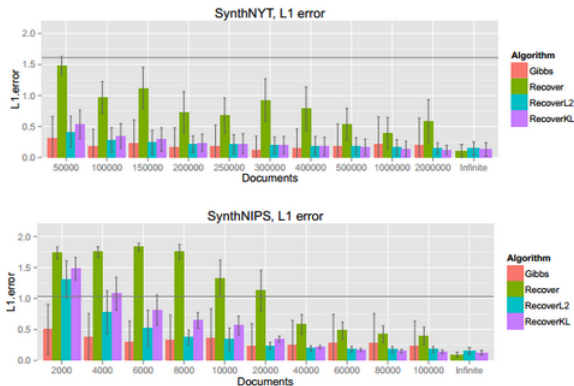


Figure 2. ℓ_1 error for learning semi-synthetic LDA models with $K = 100$ topics (**top**: based on NY Times, **bottom**: based on NIPS abstracts). The horizontal lines indicate the ℓ_1 error of K uniform distributions.

Results - Held out probability, coherence and unique words

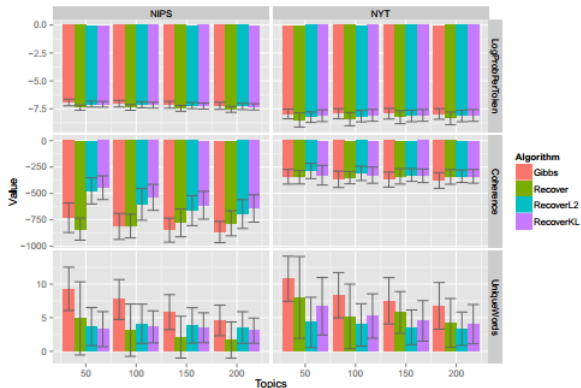


Figure 5. Held-out probability (per token) is similar for RecoverKL, RecoverL2, and Gibbs sampling. RecoverKL and RecoverL2 have better coherence, but fewer unique words than Gibbs. (Up is better for all three metrics.)



- 1 Introduction
- 2 Topic Modelling
- 3 Algorithm
- 4 Efficiently Finding Anchor Words
- 5 Topic Recovery via Bayes' Rule
- 6 Experimental Results
- 7 Conclusion**

Summary

- New algorithms for topic recovery have been presented
 - Empirical results compare to MCMC
 - Uses anchor word assumption and Bayes' rule for theoretical and empirical performance improvements
- Simple to implement, maintains provable guarantees
- Attractive feature: *Running time independent of the size of the corpus*

Future Work

- Use output of algorithms for further optimization
- Parallel implementations

Comments

- Main contribution: algorithms for topic inference with provable guarantees for running time
 - Effect: running time effectively independent of corpus size

Thanks!

Any questions please email either of us:

Vanush Vaswani

vvas9619@uni.sydney.edu.au

Kristy Hughes

khug2372@uni.sydney.edu.au