topic models: priors, stop words and languages

hanna m. wallach

university of massachusetts amherst

wallach@cs.umass.edu

Finding Needles in Haystacks



www.betaversion.org/~stefano/linotype/news/26/

- As more information becomes available, it can be harder and harder to find what we want
- We don't even always know what we want!
- Need new tools to help us organize, search and understand information

A Solution: Topic Models



Candida Hofer

- Use topic models to discover hidden topicbased patterns
- Use discovered topics to annotate the collection
- Use annotations to organize, understand, summarize, search...

Topics ↔ Words

human genome dna genetic genes sequence gene molecular sequencing map information genetics mapping project sequences

evolution evolutionary species organisms life origin biology groups phylogenetic living diversity group new two common

disease host bacteria diseases resistance bacterial new strains control infectious malaria parasite parasites united tuberculosis

computer models information data computers system network systems model parallel methods networks software new simulations

Documents \leftrightarrow **Topics**

Seeking Life's Bare (Genetic) Necessities

Haemophilus

genome 1703 genes

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

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

SCIENCE • VOL. 272 • 24 MAY 1996

"are not all that far apart," especially in comparison to the 75,000 genes in the human genome, notes Siv Andersson of Uppsala University in Sweden, who arrived at the 800 number. But coming up with a consensus answer may be more than just a genetic 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 Redundant and Related and ROM NCBI parasite-specific Genes modern genes needed genes removed removed Genes for biochemical - 4 genes -122 genes pathways. 233 genes. +22 genes ADAPTED Minimal 256 128 Mycoplasma gene set genes genes, genome 469 genes 250 genes Ancestral gene set

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

Probabilistic Modeling

- Treat data as observations that arise from a generative probabilistic process that includes hidden variables:
 - For documents, the hidden variables represent the thematic structure of the collection
- Infer the hidden structure using posterior inference:
 - What are the topics that describe this collection?
- Situate new data into the estimated model:
 - Which topics best describe the new documents?

Documents ↔ Topics ↔ Words



Probabilistic LSA

(Hofmann, 1999)



Latent Dirichlet Allocation

(Blei et al., 2003)



Dirichlet Distribution

• Distribution over K-dimensional positive vectors that sum to one (i.e., points on the probability simplex)

$$P(\boldsymbol{p} \mid \boldsymbol{\alpha}\boldsymbol{m}) = \frac{\Gamma(\sum_{k} \boldsymbol{\alpha} m_{k})}{\prod_{k} \Gamma(\boldsymbol{\alpha} m_{k})} \prod_{k} p_{k}^{\boldsymbol{\alpha} m_{k}-1}$$

- Two parameters:
 - Base measure **m** (positive vector; sums to one)
 - Concentration parameter α (positive scalar)

An Aside: The Simplex

• K-dimensional probability distributions (i.e., points on the K-1 simplex) can be plotted in (K-1)-d:



Dirichlet Parameters



Latent Dirichlet Allocation



rethinking Ida: why priors matter

hanna m. wallach, david mimno, andrew mccallum

Priors for LDA

- Almost all work on LDA uses symmetric Dirichlet priors
 - Two scalar concentration parameters: α and β
- Concentration parameters are usually set heuristically
- Some recent work on inferring optimal concentration parameter values from data (Asuncion et al., 2009)
- No rigorous study of the Dirichlet priors:
 - Base measure: asymmetric vs. symmetric
 - Treatment: optimize vs. integrate out

Topic Modeling in Practice

- Help! All my topics consist of "the, and of, to, a ..."
 - Preprocess data to remove stop words
- Now all my topics consist of "data, model, results ..."
 - Make a new corpus-specific stop word list
- Wait, but how do I choose the right number of topics T
 - Evaluate probability of held-out data for different T
- That sounds really time-consuming
 - Use a nonparametric model ...

Symmetric → Asymmetric

- Use prior over $\Theta = \{ \boldsymbol{\theta}_1, \dots, \boldsymbol{\theta}_D \}$ as a running example
- Uniform base measure \rightarrow nonuniform base measure



• Asymmetric prior: some topics more likely a priori

Predictive Distributions

- Predictive probability of topic *t* in document *d* given \mathcal{Z} $P(t \mid d, \mathcal{Z}, \alpha \boldsymbol{m}) = \int d\boldsymbol{\theta}_d P(t \mid \boldsymbol{\theta}_d) P(\boldsymbol{\theta}_d \mid \mathcal{Z}, \alpha \boldsymbol{m})$ $= \frac{N_{t \mid d} + \alpha m_t}{N_d + \alpha}$
- If t has not yet occurred in d then $P(t | d, Z, \alpha m) = m_t$
- $N_{t|d}$ is smoothed with topic-specific quantity αm_t

Handling Unknown m

- Can take a fully Bayesian approach:
 - Give **m** a Dirichlet prior: $\mathbf{m} \sim \text{Dir}(\alpha' \mathbf{u})$
 - Integrate *m* out thanks to conjugacy:

 $P(t \mid d, \mathcal{Z}, \alpha, \alpha' \boldsymbol{u}) = \int d\boldsymbol{m} P(t \mid d, \mathcal{Z}, \alpha \boldsymbol{m}) P(\boldsymbol{m} \mid \mathcal{Z}, \alpha' \boldsymbol{u})$ $= \frac{N_{t \mid d} + \alpha \frac{\hat{N}_t + \frac{\alpha'}{T}}{\sum_t \hat{N}_t + \alpha'}}{N_d + \alpha}$

An Observation

• As $\alpha' \rightarrow \infty$, the asymmetric hierarchical Dirichlet prior over Θ approaches a symmetric Dirichlet prior:

$$\frac{N_{t\mid d} + \alpha \frac{\hat{N}_t + \frac{\alpha'}{T}}{\sum_t \hat{N}_t + \alpha'}}{N_d + \alpha} \rightarrow \frac{N_{t\mid d} + \frac{\alpha}{T}}{N_d + \alpha}$$

• Symmetric Dirichlet prior is a special case of the asymmetric hierarchical Dirichlet prior

Four Combinations of Priors



Inferred Topics and Stop Words

symm. prior over Φ

- 0.080 a **field emission** an **electron** the \bigcirc
- 0.080 a the carbon and gas to an
- 0.080 the of a to and about at
- symm. 0.080 of a **surface** the with in **contact**
- 0.080 the a and to is of liquid
- 0.895 the a of to and is in
- 0.187 carbon nanotubes nanotube catalyst asymm.
- 0.043 sub is c or and n sup
- 0.061 fullerene compound fullerenes
- 0.044 material particles coating inorganic

asymm. prior over Φ

- 0.042 a **field** the **emission** and **carbon** is
- 0.042 the carbon catalyst a nanotubes
- 0.042 a the of susbtrate to material on
- 0.042 carbon single wall the nanotubes
- 0.042 the a probe tip and of to
- 1.300 the a of to and is in
- 0.257 and are of for in as such
- 0.135 a carbon material as structure nanotube
- 0.065 diameter swnt about nm than fiber swnts
- 0.029 compositions polymers polymer contain

Results: Log Probabilities



Sampled Concentration Parameters

Sampled concentration parameters from "AA"



• β' Is large compared to $\sum_{w} \hat{N}_{w}$

• Prior over Φ is effectively symmetric: "AA" \rightarrow "AS"

Intuition

- Topics are specialized distributions over words
 - Want topics to be as distinct as possible
 - Asymmetric prior over { ϕ_t } makes topics more similar to each other (and to the corpus word frequencies)
 - Want a symmetric prior to preserve topic "distinctness"
- Still have to account for power-law word usage:
 - Asymmetric prior over { θ_d } means some topics (e.g., "the, a, of, to ...") can be used more often than others

Conclusions

- Careful thinking about priors can yield new insights
 - e.g., priors and stop word handling are related
- For LDA the choice of prior is surprisingly important:
 - Asymmetric prior for document-specific topic distributions
 - Symmetric prior for topic-specific word distributions
- Rethinking priors for LDA facilitates new topic models
 - e.g., polylingual topic model ...

NESCAI 2010

- April 16-18 @ UMass Amherst
- http://nescai.cs.umass.edu/cfp.php



questions?