

# EFFICIENT POSTERIOR SAMPLING IN TOPIC MODELS FOR TEXT

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# LECTURE OVERVIEW

- Textual data
- Topic models
- Efficient posterior sampling in topic models

# TEXTUAL DATA

- **Digitalization**: text is becoming an important data source.
- The web, PDF documents (legal, political, medical, etc)
- Unstructured (not tables), but structured (by the rules of language).
- **Big data**. 100K, 1M or even more documents in a data set.
- Lots of **pre-processing** to get the data in usable form for statistical analysis.

# TEXT APPLICATIONS

- **Language models** (predict the next word on smartphone keyboard)
- **Machine translation** (Google translate)
- **Document classification** (Shakespeare? Spam and blog filters. harmful EULA? pos/neg financial statement?)
- **Information retrieval** (Google search)
- **Sentiment analysis** (positive/negative sentiment in tweets)
- **Part-of-speech tagging** (classify the grammatical category of words in a sentence)
- **Prediction models** based on text. Predicting financial turbulence from econ media. Media-Financial markets feedback.

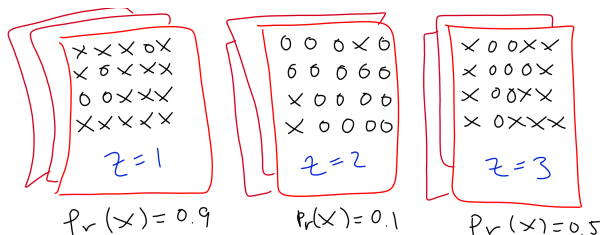
# TOPIC MODELS

- **Latent Dirichlet Allocation.**
- **Probabilistic generative** models for text documents, but also useful in other settings.
- **Very popular** model in applications and research. > 12000 Google scholar citations in 11 years.
- Topic models summarizes documents into low-dimensional **topics**. **Dimension reduction**. “PCA for discrete data”.
- **Many extensions** in recent years.

# MIXTURE OF UNIGRAMS

- **Mixture of unigrams:**

1. Draw a **topic**  $z_d$  for the  $d$ th document from a topic distribution  $\theta = (\theta_1, \dots, \theta_K)$ .
2. Conditional on the drawn topic  $z_d$  draw words from a **word distribution** for that topic.



# TOPIC MODELS

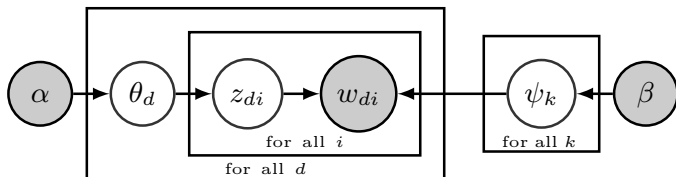
- Mixture of unigrams: each document belong to exactly one topic.
- **Topic models** are **mixed-membership models**: each document can belong to **several topics simultaneously**.
- More general than text. Individuals' behavior is determined by a mixture of influences.

# GENERATING A CORPUS FROM A TOPIC MODEL

- Assume that we have:
  - A fixed vocabulary  $V$
  - $D$  documents
  - $N$  words in each document
  - $K$  topics
- 1. For each **topic** ( $k = 1, \dots, K$ ):
  - A. Draw a distribution over the words  $\phi_k \sim \text{Dir}(\eta, \eta, \dots, \eta)$
- 2. For each **document** ( $d = 1, \dots, D$ ):
  - A. Draw a vector of topic proportions  $\theta_d \sim \text{Dir}(\alpha_1, \dots, \alpha_K)$
  - B. For each **word** ( $i = 1, \dots, N$ ):
    - I. Draw a topic assignment  $z_{di} \sim \text{Multinomial}(\theta_d)$
    - II. Draw a word  $w_{di} \sim \text{Multinomial}(\phi_{z_{di}})$



# GENERATING A CORPUS FROM A TOPIC MODEL



# EXAMPLE - SIMULATION FROM TWO TOPICS

Topic	Word distr.	probability	dna	gene	data	distribution
1	$\phi_1$	0.5	0.1	0.0	0.2	0.2
2	$\phi_2$	0.0	0.5	0.4	0.1	0.0

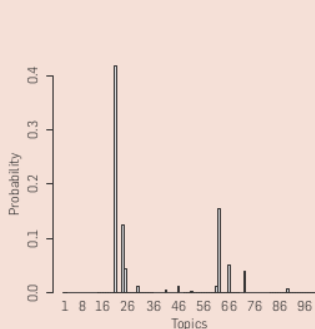
Doc 1	$\theta_1 = (0.2, 0.8)$				
	Word 1:	Topic=2	Word='gene'		
	Word 2:	Topic=2	Word='gene'		
	Word 3:	Topic=1	Word='data'		

Doc 2	$\theta_2 = (0.9, 0.1)$				
	Word 1:	Topic=1	Word='probability'		
	Word 2:	Topic=1	Word='data'		
	Word 3:	Topic=1	Word='probability'		

Doc 3	$\theta_2 = (0.5, 0.5)$				
⋮	⋮	⋮	⋮	⋮	⋮

# EXAMPLE FROM SCIENCE (BLEI, REVIEW PAPER)

Figure 2. Real inference with LDA. We fit a 100-topic LDA model to 17,000 articles from the journal *Science*. At left are the inferred topic proportions for the example article in Figure 1. At right are the top 15 most frequent words from the most frequent topics found in this article.



<b>"Genetics"</b>	<b>"Evolution"</b>	<b>"Disease"</b>	<b>"Computers"</b>
human	evolution	disease	computer
genome	evolutionary	host	models
dna	species	bacteria	information
genetic	organisms	diseases	data
genes	life	resistance	computers
sequence	origin	bacterial	system
gene	biology	new	network
molecular	groups	strains	systems
sequencing	phylogenetic	control	model
map	living	infectious	parallel
information	diversity	malaria	methods
genetics	group	parasite	networks
mapping	new	parasites	software
project	two	united	new
sequences	common	tuberculosis	simulations

# GIBBS SAMPLER

- **Posterior distribution**

$$p(\mathbf{z}, \Theta, \Phi | \mathbf{w}) \propto p(\mathbf{z}, \Theta, \Phi | \mathbf{w}) \cdot p(\mathbf{z}, \Theta, \Phi)$$

- Integrating out (**collapsing**)  $\Theta$  and  $\Phi$ :

$$p(\mathbf{z} | \mathbf{w}) = \int \int p(\mathbf{z}, \Theta, \Phi | \mathbf{w}) \cdot p(\mathbf{z}, \Theta, \Phi) d\Phi d\Theta$$

will result in the following **Gibbs sampler** for the  $\mathbf{z}$ 's

$$p(z_{di} = k | w_{di} = w, \mathbf{z}_{-di}) = \underbrace{\frac{n_{k,w}^{-di} + \beta}{n_{k,\cdot}^{-di} + V\beta}}_{\text{type-topic } (\Phi)} \cdot \underbrace{(n_{k,d}^{-di} + \alpha)}_{\text{topic-doc } (\Theta)}.$$

- The rows of  $\Phi | \mathbf{z}$  and  $\Theta | \mathbf{z}$  are Dirichlet.
- The inferred topic proportions  $\theta_d$  are useful summaries of document content that can be used as covariates in regression/classification.
- Example: predicting financial events from news articles.

# (NAIVE) ALGORITHM

```
# Initialization
```

```
Sample all topic indicators randomly
```

```
Calculate  $n^{(w)}$  and  $n^{(d)}$  # topic-word and topic-document  
# matrices with counts
```

```
# Gibbs sampler
```

```
for each gibbs iteration do
```

```
  for each token  $w_i$  do
```

```
    remove  $z_i$  from  $n^{(w)}$  and  $n^{(d)}$ 
```

```
    for each  $k$  in 1 to  $K$  do
```

$$\text{prob}_k[k] = \frac{n_{k,v_i}^{(w)} + \beta}{n_{k,\cdot}^{(w)} + V\beta} \cdot (n_{k,d_i}^{(d)} + \alpha)$$

```
    end for
```

```
     $z_i \leftarrow$  draw multinomial(prob_k)
```

```
    add  $z_i$  to  $n^{(w)}$  and  $n^{(d)}$ 
```

```
  end for
```

```
end for
```

# SPEEDING UP MCMC FOR TOPIC MODELS

- **Gibbs sampling** from  $p(z_{di} = k | w_{di} = w, z_{-di})$  is **slow** since it runs serially over all tokens in the corpus.
- Example: PubMed abstracts. 10% of the data: **78.5M tokens** from **820K docs**.
- Lesson: with a huge number of parameters, everything matters ( $\log(x)$  is costly ....).
- Needed: **Efficient data structures, sparsity, efficient search, efficient sort** etc.

# SPEEDING UP MCMC FOR TOPIC MODELS

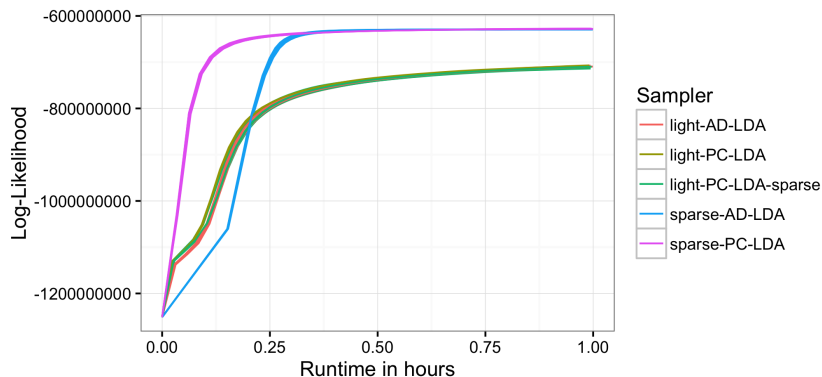
1. **Efficient sparse data structures.** Lots of binaries. Only store non-zero positions.
2. **Partially collapsing.** Integrate out  $\Theta$ , but not  $\Phi$ . Documents are now conditionally independent:  $p(\mathbf{z}_1, \dots, \mathbf{z}_D | \Phi, \mathbf{w}) = \prod_{d=1}^D p(\mathbf{z}_d | \Phi, \mathbf{w})$ . **Document-parallel sampling** of the  $\mathbf{z}$ 's. Very little efficiency loss from not integrating out  $\Phi$  [1]
3. **Exploit sparsity.** Sampling each  $z_{id}$  has complexity  $O(K)$ . **Walker's Alias**: fast way to repeatedly sample from dense discrete distributions that do not change much from iteration to iteration.

$$p(z_{di} = k | \cdot) = \underbrace{n_{k,d}^{-di} \frac{n_{k,w}^{-di} + \beta}{n_{k,\cdot}^{-di} + V\beta}}_{\text{Sampling in } O(K_d)} + \underbrace{\alpha \frac{n_{k,w}^{-di} + \beta}{n_{k,\cdot}^{-di} + V\beta}}_{\text{Alias sampling in amortized } O(1)}$$

MH-step needed for the full collapsed sampler. Simple Gibbs for partially collapsed.

# PUBMED EXAMPLE

- Abstracts from medical articles.
- Here we use 10% of the data. 78.5M tokens from 820K docs.





# TOPIC MODEL AS BUILDING BLOCK IN OTHER MODELS

- Predicting the location of bugs in computer code [2]. Ericsson.
- New **multi-class** (code component) prediction model for high-dimensional data.
- **Diagonal Orthant Multinomial Probit** [3] instead of the usual multinomial logit/probit. No reference category.
- Prediction machine: Bug report  $\Rightarrow$   $\text{Pr}(\text{Component})$ .
- **DOLDA** - Diagonal Orthant Latent Dirichlet Allocation
- **Interpretable predictions** via semantical topics and **aggressive horseshoe regularization**.

# PREDICTING BUG LOCATION FROM BUG REPORTS

```
#
def Network(networkInputs):
```

```
# CODE
# MORE CODE
# TOO MUCH CODE
```

```
return(networkOutputs)
```

```
#
def UI(UIinputs):
```

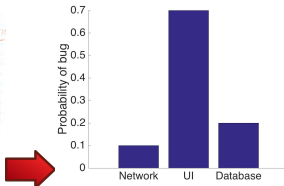
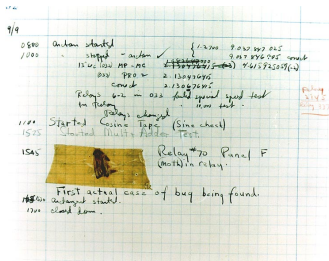
```
# CODE
# MORE CODE
# TOO MUCH CODE
```

```
return(UIOutputs)
```

```
#
def Database(DBinputs):
```

```
# CODE
# MORE CODE
# TOO MUCH CODE
```

```
return(DBOutputs)
```



# EXAMPLE DATA

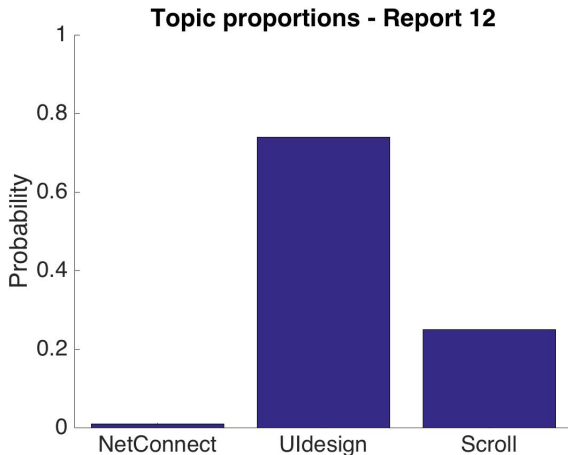
Dataset	No. Bug reports	No. components	Vocabulary size
Mozilla	15,000	118	3505
Eclipse	15,000	49	3367
Telecom	9,778	26	5286

# TOPICS $\phi_k$

- Automatically summarize a bug report by topics.

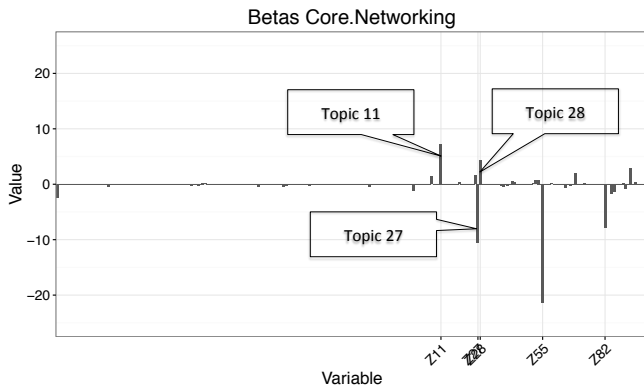
Topic	Topic label	Top 10 words in topic
11	HTTP	proxy server http network connection request connect error www host
27	Layout	div style px background color border css width height element html
28	Connection Headers	http cache accept en public local-host gmt max modified alive
55	Search	search google bar results box type find engine enter text
82	Scrolling	scroll page scrolling mouse scrollbar bar left bottom click content

# TOPIC PROPORTIONS $\theta_d$



# IDEAL: FEW TOPICS FOR EACH COMPONENT

- Horseshoe regularization prior.



# INTERPRETABLE PREDICTIONS

- Our model: **DOLDA** - Diagonal Orthant Latent Dirichlet Allocation. Supervised LDA. Topics are directly related to components.
- System: *"I am very certain that component UI contains the bug because report talks a lot about Uldesign and Scroll and very little about NetConnect."*
- System: *"I am very uncertain where the bug is because bug report contains a jumble of topics. Please ask human."*

# NO INTERPRETATION VS ACCURACY TRADE-OFF

Dataset	DOLDA-normal	DOLDA-horseshoe	StackingLDA	LDA-KL
Mozilla	46%	45%	39%	26%
Eclipse	61%	61%	55%	37%
Telecom	72%	71%	75%	41%





M. Magnusson, L. Jonsson, M. Villani, and D. Broman, “Parallelizing lda using partially collapsed gibbs sampling,” *arXiv preprint arXiv:1506.03784*, 2015.



L. Jonsson, M. Magnusson, D. Broman, K. Sandahl, M. Villani, and S. Eldh, “Automatic localization of bugs to faulty components in large scale software systems using bayesian classification,” in *IEEE Conference on Software Quality and Reliability (QRS-16)*, 2016.



J. Johndrow, D. B. Dunson, and K. Lum, “Diagonal orthant multinomial probit models,” in *AISTATS*, pp. 29–38, 2013.