

#### COMP90042 LECTURE 15

# TOPIC MODELS

### INTRODUCTION

- ► A librarian team just digitalised thousands of documents. Now they need to organise them.
- Twitter wants to find which topics are trending today.
   (half a billion tweets per day)
- A company wants to make sense of thousands of reports from its employees.
- Maybe we can use text classification?
  - Get some annotated data, train a classifier, predict classes for unseen texts, report.

### INTRODUCTION

- A librarian team just digitalised thousands of documents. Now they need to organise them.
  - What about documents with multiple classes?
- Twitter wants to find which topics are trending today.
   (half a billion tweets per day)
  - What about new classes?
- A company wants to make sense of thousands of reports from its employees.
  - ▶ What **are** classes?

### PROBLEMS WITH TEXT CLASSIFICATION

- Simple solutions based on text classification not feasible.
  - Documents can have multiple classes (multidisciplinary books, for instance).
  - Lack of annotated data.
  - Sometimes we don't even know which classes we expect from the data or we are interested into. Remember unsupervised information extraction (OpenIE)?

### **TOPICS**

- We need a concept which is more open-ended than "classes": **topics**.
- A topic can have a specific semantic interpretation but does not have a label.
- Additionally, documents can have multiple topics.
- This concept enables us to perform open-ended, unsupervised learning on documents.
  - ► The standard algorithm for this is called Latent Dirichlet Allocation (LDA). We will derive it in this lecture.

### TOPICS - EXAMPLE

"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."

### TOPICS - EXAMPLE

- "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."
- ► Topic 1: "genes", "genome", "genomes"
- ▶ Topic 2: "organism", "survive", "life", "organisms"
- ► Topic 3: "computer"
- ▶ **Key assumption:** words within a document have a single topic.

### TOPICS - EXAMPLE

- ▶ Topic 1: "genes", "genome", "genomes"
- ▶ Topic 2: "organism", "survive", "life", "organisms"
- ► Topic 3: "computer"
- We can assign topics to documents based on topic frequency, for instance.
  - Document 1: topics 2 and 3
  - Document 2: topics 1 and 3
- ▶ What does topic "1" mean?
  - Need to inspect the words and manually assign semantics.
  - ▶ We will revisit this later.

- "How many genes does an organism need to survive?"
- How/0 many/0 genes/1 does/0 an/0 organism/2 need/0 to/0 survive/2 ?/0
- First idea: unsupervised HMMs
  - Split each document into sentences.
  - Assume, for instance, 100 topics. Initialise, run EM.
- Assign topics based on frequency.
  - Problem: this will likely assign the same topics to every document.
  - Limitation: a word's topic is independent of the document.

- Second idea: per-document unsupervised HMMs.
  - ► The topic distribution now varies for each document, which is exactly we are aiming for.
  - But this simple approach will not account for how topics relate across documents. Topic "4" in document "1" might be the same as topic "7" in document "5". We need to **tie** the parameters across all documents.

- Let's formalise that:
  - P(w = gene, t = 1 | d = 7) =
  - P(w = gene|t = 1, d = 7) \* P(t = 1|d = 7)
- ► Two key simplifications:
- Emission probabilities are shared across documents
  - P(w = gene|t = 1, d = 7) = P(w = gene|t = 1)
- Topic of word does not depend on previous words' topics.
  Only depends on the document.
  - P(t = 1|d = 7) becomes a single parameter.

- P(w = gene, t = 1 | d = 7) =
  - $P(w = \text{gene}|t = 1, \beta_1) * P(t = 1|\theta_7)$
- Parameters are now:
  - $\triangleright$   $\beta$  (one per **topic**): the distribution of words given a topic
  - $\theta$  (one per **document**): the distribution of topics given a document
- ▶ We can use EM to train this. Given a topic initialisation:
  - $\beta$ : count (expected) word-topic frequencies (in the whole corpus) and normalise
  - $\bullet$ : count (expected) topic-document frequencies and normalise

### EM FOR TOPIC MODELLING

E-step

$$P(t = 1 | w = \text{gene}, d = 7, \beta_1, \theta_7) = \frac{P(w = \text{gene}, t = 1 | d = 7, \beta_1, \theta_7)}{\sum_k P(w = \text{gene}, t = k | d = 7, \beta_k, \theta_7)}$$

M-step

$$\theta_7^1 = \frac{\sum_{i \in d} P(t=1|w_i,\beta_1,\theta_7)}{\sum_k \sum_{i \in d} P(t=k|w_i,\beta_k,\theta_7)} \approx \frac{\text{# tokens in } d \text{ with topic 1}}{\text{# tokens in } d}$$

$$\beta_1^w = \sum_{d \in D} \frac{\sum_{i: w_i = w} P(t=1|w_i = w, \beta_1, \theta_d)}{\sum_{v \in V} \sum_{i: w_i = v} P(t=k|w_i = v, \beta_1, \theta_7)}$$

\*  $\approx$   $\frac{\text{# tokens with type } w \text{ and with topic 1}}{\text{# tokens with topic 1}}$ 

- ▶ We are almost there! The previous model is very close to Latent Dirichlet Allocation.
- It is missing a key component that we saw in other probabilistic models before: **smoothing**.
- Let's do add-k smoothing.

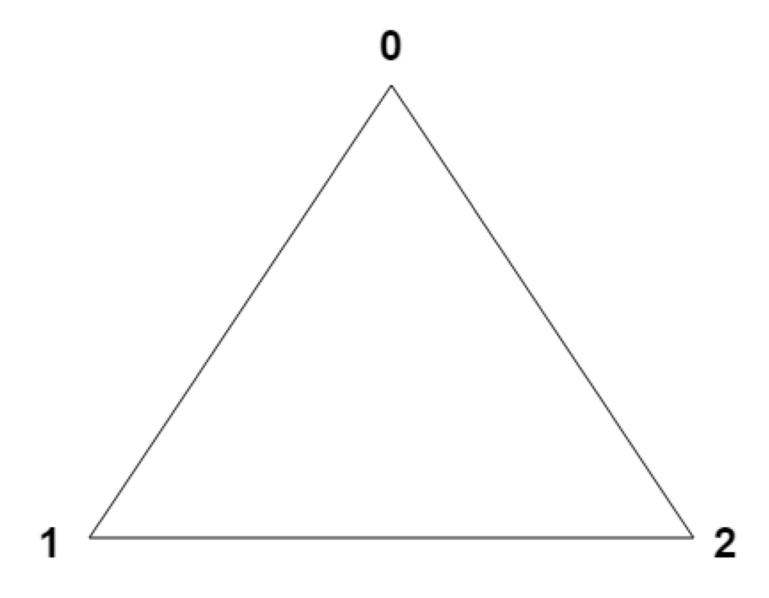
### **ADD-K SMOOTHING**

- E-step is the same
- M-step

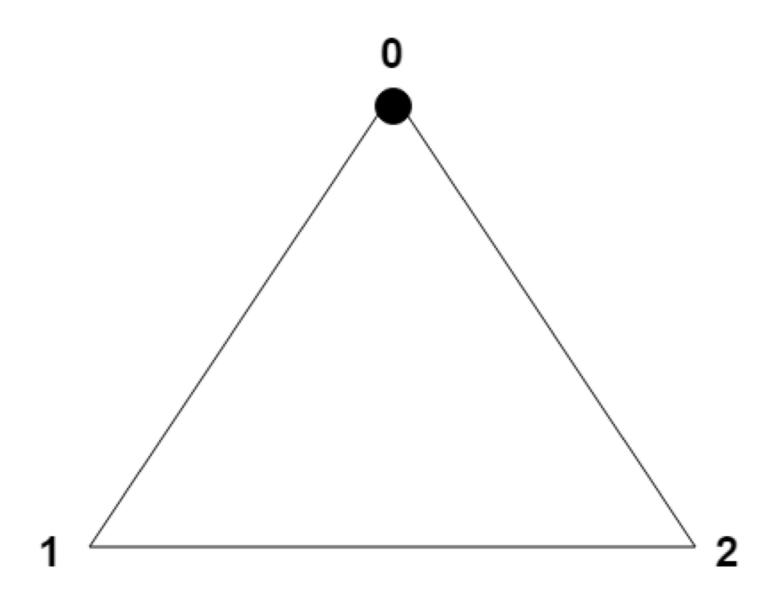
$$\bullet \ \theta_7^1 = \frac{\alpha + \sum_{i \in d} P(t=1|w_i, \beta_1, \theta_7)}{\sum_{k} (\alpha + \sum_{i \in d} P(t=k|w_i, \beta_k, \theta_7))}$$

$$\beta_1^w = \sum_{d \in D} \frac{\eta + \sum_{i:w_i = w} P(t=1|w_i = w, \beta_1, \theta_d)}{\sum_{v \in V} (\eta + \sum_{i:w_i = v} P(t=k|w_i = v, \beta_1, \theta_7))}$$

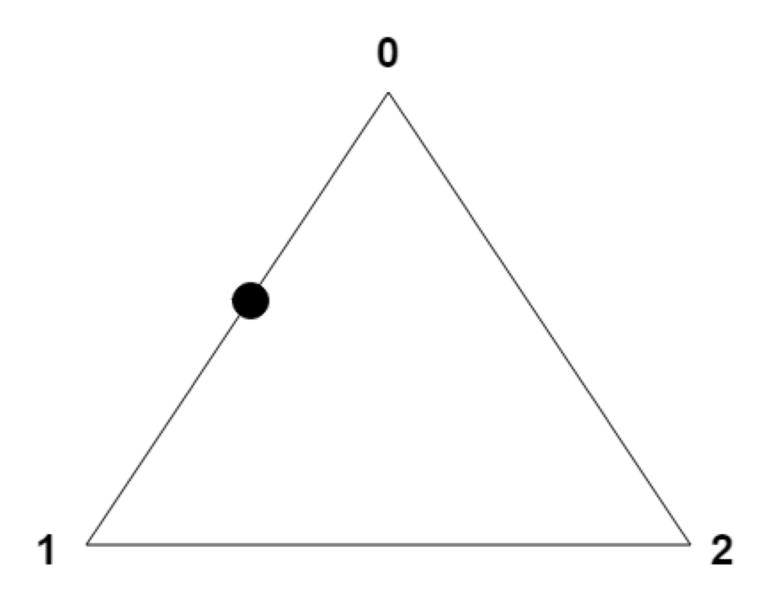
- Add-k smoothing can be interpreted as having a **prior distribution** over the parameters  $\theta$  and  $\beta$ 
  - $P(\theta|\alpha) = Dirichlet(\alpha + 1)$
  - $P(\beta|\eta) = Dirichlet(\eta + 1)$



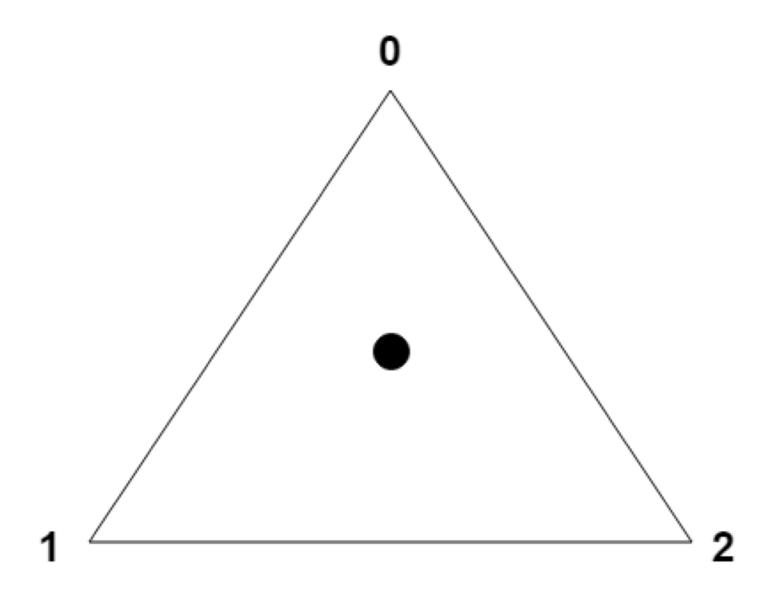
► Each point in the triangle is a probability distribution over 3 topics



$$P(0) = 1, P(1) = 0, P(2) = 0$$

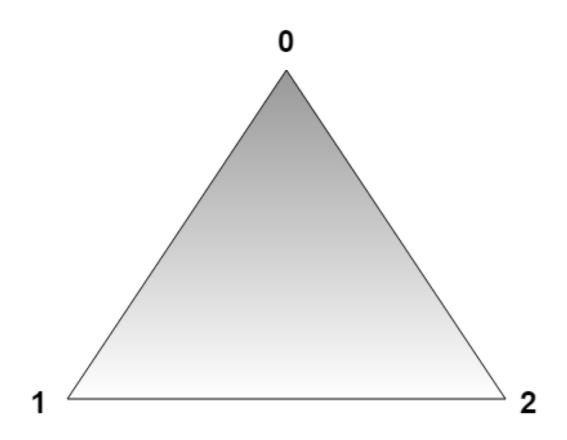


$$P(0) = 0.5, P(1) = 0.5, P(2) = 0$$



$$P(0) = 0.333, P(1) = 0.333, P(2) = 0.333$$

### THE DIRICHLET DISTRIBUTION



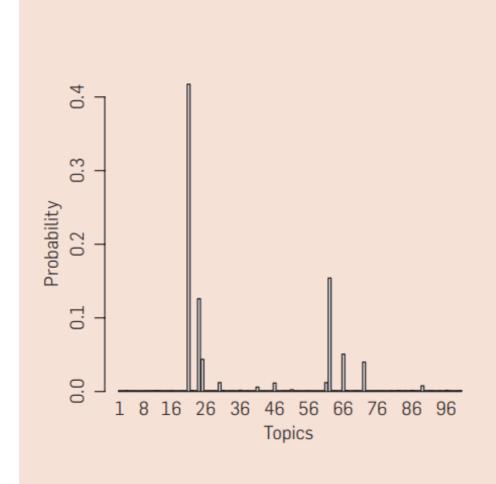
- ▶ The Dirichlet is a distribution over a probability simplex.
  - $\triangleright$   $\alpha$  defines the spread
  - Symmetric:  $\alpha$  is a scalar
  - Asymmetric: one  $\alpha$  per topic

### LATENT DIRICHLET ALLOCATION

- ► The model we saw before is essentially LDA with symmetric Dirichlet over topics and words
  - Parameters can be estimated via EM
- ► In practice:
- Use asymmetric Dirichlet
- Instead of finding the maximum value for  $\theta$  and  $\beta$ , estimate the **posterior** distribution over the parameters
  - Harder to do but prevents overfitting
  - ► Usually done via Gibbs sampling or variational inference: we will not see these techniques here... ⊗

- Intrinsic: perplexity.
  - ► The model defines a distribution over each word, same as a language model.
  - We can split the corpus into a training and test set and evaluate perplexity on the test set.
- Extrinsic is harder.
  - If the topic model is used as a tool for an end task (information retrieval, machine translation, etc.), calculate the end task metric.
  - Otherwise some human intervention is required. Interpretability is key.

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.



| "Genetics"  | "Evolution"  | "E  |
|-------------|--------------|-----|
| human       | evolution    | (   |
| genome      | evolutionary |     |
| dna         | species      | k   |
| genetic     | organisms    | C   |
| genes       | life         | re  |
| sequence    | origin       | b   |
| gene        | biology      |     |
| molecular   | groups       |     |
| sequencing  | phylogenetic |     |
| map         | living       | in  |
| information | diversity    | 1   |
| genetics    | group        | ķ   |
| mapping     | new          | р   |
| project     | two          |     |
| sequences   | common       | tul |
|             |              |     |

| "Disease"    | "Computers  |
|--------------|-------------|
| disease      | computer    |
| host         | models      |
| bacteria     | information |
| diseases     | data        |
| resistance   | computers   |
| bacterial    | system      |
| new          | network     |
| strains      | systems     |
| control      | model       |
| infectious   | parallel    |
| malaria      | methods     |
| parasite     | networks    |
| parasites    | software    |
| united       | new         |
| tuberculosis | simulations |
|              |             |

Blei (2012)

Figure 3. A topic model fit to the *Yale Law Journal*. Here, there are 20 topics (the top eight are plotted). Each topic is illustrated with its topmost frequent words. Each word's position along the *x*-axis denotes its specificity to the documents. For example "estate" in the first topic is more specific than "tax."

| 4             | 10          | 3                      | 13             |
|---------------|-------------|------------------------|----------------|
| tax           | labor       | women                  | contract       |
| income        | workers     | sexual                 | liability      |
| taxation      | employees   | men                    | parties        |
| axes          | union       | sex                    | contracts      |
| revenue       |             | child                  |                |
|               | employer    |                        | party          |
| estate        | employers   | family                 | creditors      |
| subsidies     | employment  | children               | agreement      |
| exemption     | work        | gender                 | breach         |
| organizations | employee    | woman                  | contractual    |
| year          | job         | marriage               | terms          |
| treasury      | bargaining  | discrimination         | bargaining     |
| consumption   | unions      | male                   | contracting    |
| taxpayers     | worker      | social                 | debt           |
| earnings      | collective  | female                 | exchange       |
| funds         | industrial  | parents                | limited        |
| 6             |             | 1                      | 10             |
| 0             | 15          | 1                      | 16             |
| jury          | speech      | firms                  | constitutional |
| trial         | free        | price                  | political      |
| crime         | amendment   | corporate              | constitution   |
| defendant     | freedom     | firm                   | government     |
| defendants    | expression  | value                  | justice        |
| sentencing    | protected   | market                 | amendment      |
| judges        | culture     | cost                   | history        |
| punishment    | context     | capital                | people         |
| judge         | equality    | shareholders           | legislative    |
| crimes        | values      | stock                  | opinion        |
| evidence      | conduct     |                        | fourteenth     |
| sentence      | ideas       | insurance<br>efficient | article        |
|               | information |                        |                |
| jurors        |             | assets                 | majority       |
| offense       | protect     | offer                  | citizens       |
| guilty        | content     | share                  | republican     |

Blei (2012)

- Many other visualisation options available
  - Word clouds
  - Word graphs
- Labelling techniques
  - Distant supervision (Wikipedia, knowledge bases, etc.)
  - Article names, pictures.
- Combination of all above
  - Goal: enhance interpretability

### LDA EXTENSIONS

- LDA is the standard method for topic modelling. It has been vastly studied in academia and many extensions were proposed.
- LDA-HMM: nearby words have similar topics
- Hierarchical LDA: assumes a topic hierarchy
- Correlated LDA: some topics are more similar to others ("football" and "rugby" vs. "football" and "genetics")
- Dynamic LDA: topics change over time, need to know document ordering (from timestamps, for instance)
- Non-parametric LDA: does not assume a fixed number of topics (harder to train)

### **APPLICATIONS**

- Information retrieval: useful for query expansion.
- Analysis of historical documents (newspapers, books).
- Making sense of scientific publications: emergence of new fields and multidisciplinary ones.
- Literary analysis: stylometry, comparative literature.
- Computational social science: text on social media (Twitter), stance detection.
- Translation: multilingual topic models for mining parallel data.

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#### A FINAL WORD

- ► Topic models aim at making sense of large collections of documents, combining two ideas:
  - Unsupervised learning (clustering)
  - Documents can have multiple topics
- ► LDA is the standard method
  - Can use EM to train via MAP
  - In practice, estimate posteriors using sampling or other techniques
- Perplexity can give evidence of performance but ultimate goal in clustering is interpretability

#### ADDITIONAL READING

- David Blei's ACM paper
   (<a href="http://www.cs.columbia.edu/~blei/papers/Blei2012.pdf">http://www.cs.columbia.edu/~blei/papers/Blei2012.pdf</a>)
- Optional) "Applications of Topic Models"
  - https://mimno.infosci.cornell.edu/papers/2017\_fntir\_tm\_ap plications.pdf