LATENT VARIABLES AND NATURAL LANGUAGE PROCESSING

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LATENT VARIABLES AND NATURAL LANGUAGE PROCESSING

LEARNING OBJECTIVES

- Understand what *latent* variables are
- Understand the uses of *latent variables* in language processing
- Use the *word2vec* and *LDA* algorithms of genism

COURSE

PRE-WORK

PRE-WORK REVIEW

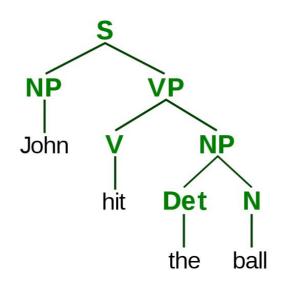
- Install gensim with pip install gensim
- Recall and apply unsupervised learning techniques
- Recall probability distributions, specifically discrete multinomial distributions
- Recall NLP essentials, including experience with spacy

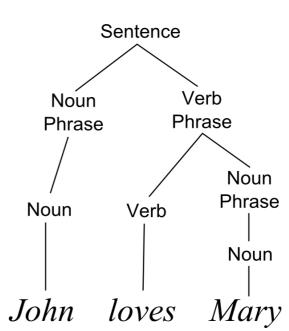
LATENT VARIABLE MODELS vs TRADITIONAL NLP

- This lesson will continue on natural language processing with an emphasis on *latent variables models*.
- Mining and Refining data is a key part of the data science workflow.
- During the previous lessons, we saw many techniques for mining the data, including preprocessing, building linguistic rules to uncover patterns, and creating classifiers from unstructured data.
- In this class, we'll continue with methods to Refine our understanding of the text by attempting to uncover structure or organization in the text.

- Many advances in NLP are based on using data to learn rules of grammar and language. We saw these tools in our last class.
 - Tokenization
 - Stemming or lemmatization
 - Parsing and tagging
- Each of these are based on a classical or theoretical understanding of language.

- Tokenization:
 - \rightarrow John hit the ball \rightarrow [John, hit, the, ball]
 - Where did you go → [Where, did, you, go]
- Stemming or lemmatization: shouted \rightarrow shout, better \rightarrow good
- Parsing and tagging:





- Latent variable models are different in that they try to understand language based on **how** the words are used.
- For example, instead of learning that 'bad' and 'badly' are related because they share the same root, we'll determine that they are related because they are often used in the same way often or near the same words.
- We'll use *unsupervised* techniques (discovering patterns or structure) to extract the information.

Traditional NLP Models

Focused on theoretical understanding of language

Tries to learn the rules of a particular language

Preprogrammed set of rules

Latent Variable Models

Focused on how the language is actually used in practice

Infers meaning from how words are used together

Uses unsupervised learning to discover patterns or structure

Traditional NLP Models

'bad' and 'badly' are related because they share a common root.

'Python' and 'C++' are both programming languages because they are often a noun preceded by the verb 'program' or 'code'.

Latent Variable Models

'bad' and 'badly' are related because they are used the same way or near the same words.

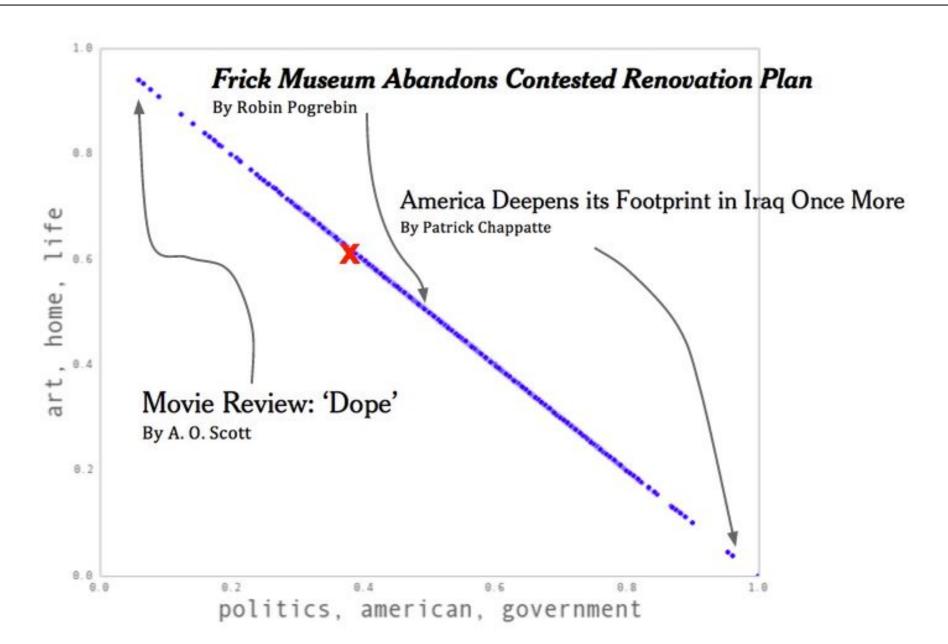
'Python' and 'C++' are both programming languages because they are often used in the same context.

INTRODUCTION

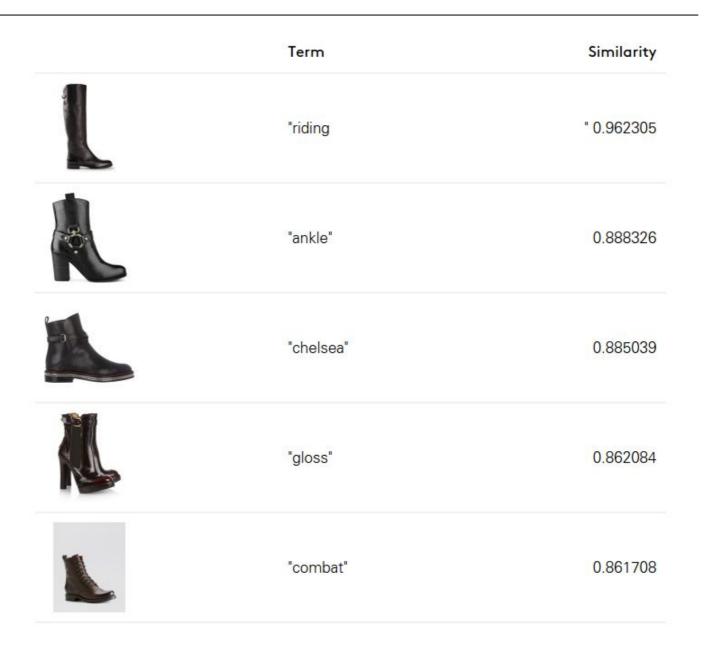
- Latent variable models are models in which we assume the data we are observing has some **hidden**, **underlying structure** that we can't see, and which we'd like to learn.
- These hidden, underlying structures are the *latent* (i.e. hidden) variables we want our model to understand.
- Text processing is a common application of latent variables.
- The goal is to infer a model that would produce our text.

- While language (in the classical sense) is defined by a set of pre-structured grammar rules and vocab, we often break those rules and create new words (e.g. selfie).
- Instead of attempting to train our model on the rules of proper grammar, we'll ignore grammar and seek to uncover alternate hidden structures.

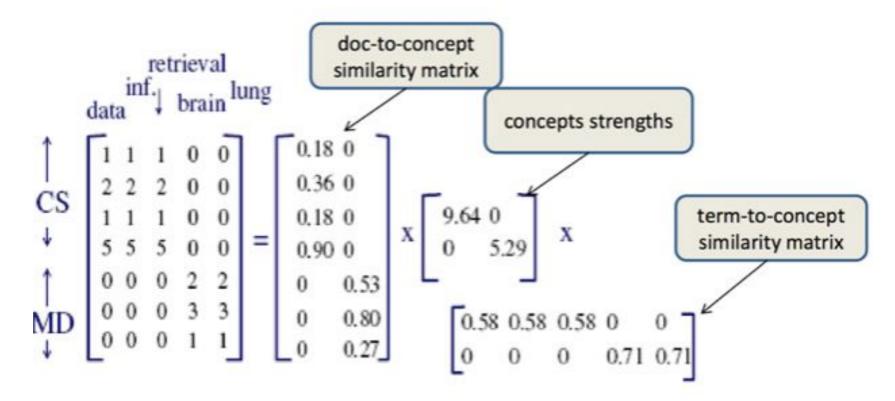
- Latent variable techniques are often used for recommending news articles or mining large troves of data to find commonalities.
- Topic modeling, a method we'll cover today, is used in the NY times recommendation engine.
- The New York Times attempts to map their articles to a latent space of topics using the content of the article.



Lyst, an online fashion retailer, uses latent representations of clothing descriptions to find similar clothing.



• Our previous 'representation' of a set of text documents (articles) for classification was a matrix with one row per document and one column per word (or n-gram).



- While this sums up most of the information, it does drop a few things, mostly structure and order.
- Additionally, many of the columns may be correlated.

- For example, an article that contains the word 'IPO' is likely to contain the word 'stock' or 'NASDAQ'.
- Therefore, those columns are repetitive and likely to represent the same concept or idea.
- For classification, we may only care that there are finance-related words.

- One way to deal with this is through regularization L1/Lasso regularization tends to remove repetitive features by bringing their learned coefficients to 0.
- Another is to perform *dimensionality reduction*, where we first identify the correlated columns and the replace them with a column that represents the concept they have in common.
- For instance, we could replace 'IPO', 'stocks', and 'NASDAQ' with a single column 'HasFinancialWords' column.

- There are many techniques to do this automatically and mostly follow a very similar approach.
 - a. Identify correlated columns.
 - b. Replace them with a new column that encapsulates the others.

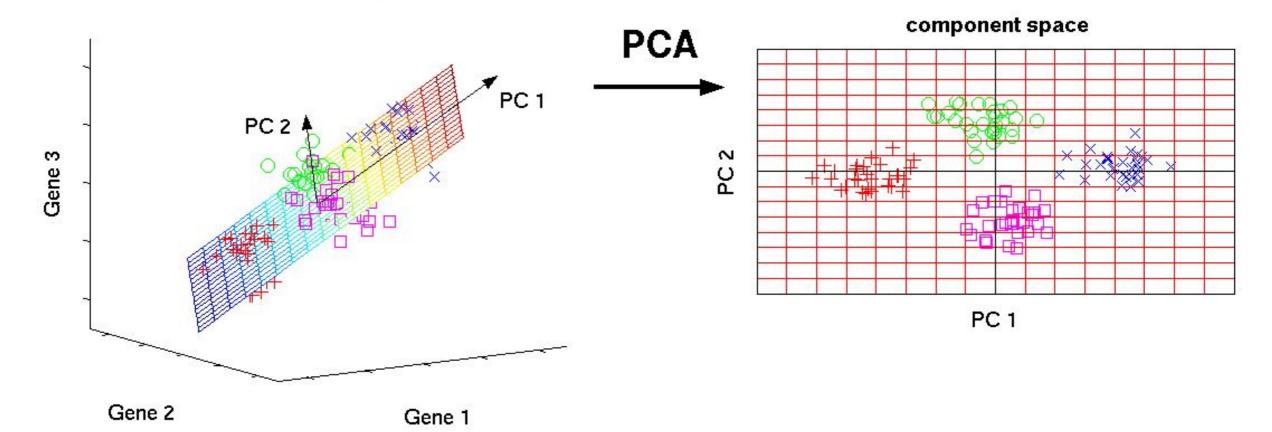
Doc #	Car	Truck	Van	Dog	Doc #	Vehicle	Dog
6344	1	1	1	0	6344	1	О
6345	0	1	1	1	6345	1	1
6346	1	1	1	0	6346	1	О

- The techniques vary in how they define correlation and how much of the relationship between the original and new columns you need to save.
- Dimensionality techniques can vary between *linear* and *non-linear*.

- There are many techniques build into scikit-learn.
- One of the most common is **Principal Component Analysis** (**PCA**).
- PCA like techniques, when applied to text data, is sometimes known as <u>Latent Semantic Indexing (LSI)</u>.
- http://scikit-learn.org/stable/modules/decomposition.html#truncated-s ingular-value-decomposition-and-latent-semantic-analysis

• PCA helps reduce the feature space into fewer dimensions.

original data space



- Mixture models (specifically **LDA** or **Latent Dirichlet Allocation**) take this concept further and generate more structure around the documents.
- Instead of just replacing correlated columns, we create clusters of common words and generate probability distributions to explicitly state how related words are.

- To understand this better, let's imagine that we generate text by:
 - a. Start writing a document
 - i. Choose a topic (sports, news, science).
 - ii. Choose a random word from that topic.
 - iii. Repeat.
 - b. Repeat for the next document.

- This 'model' of text is assuming that each document is some *mixture* of topics.
- It may be mostly science but may contain some business information.
- The *latent* structure we want to uncover are the topics (or concepts) that generate that text.

- Latent Dirichlet Allocation is a model that assumes this is the way text is generated and then attempts to learn two things:
 - a. The word distribution of each topic
 - b. The topic distribution of each document.

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

Topic proportions and assignments

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.

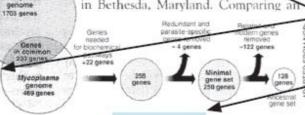
Although the numbers don't match precisely, those predictions

* Genome Mapping and Sequencing, Cold Spring Harbor, New York,

May 8 to 12.

"are not all that far apart," especially in comparison to the 75,000 genes in the human genome, notes Siv Andersson of the sala University in Sweden, the 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

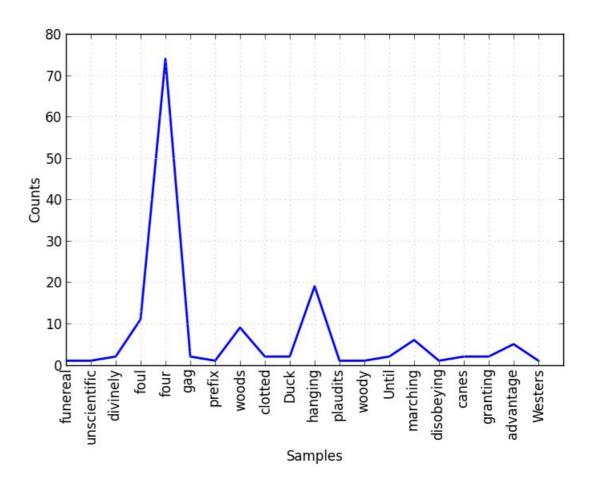
Arcady Mushegian, a computational molecular biologist at the National Center for Biotechnology Information (NCBI) in Bethesda, Maryland. Comparing an



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

SCIENCE • VOL. 272 • 24 MAY 1996

The word distribution is a multinomial distribution of each topic representing what words are most likely from that topic.



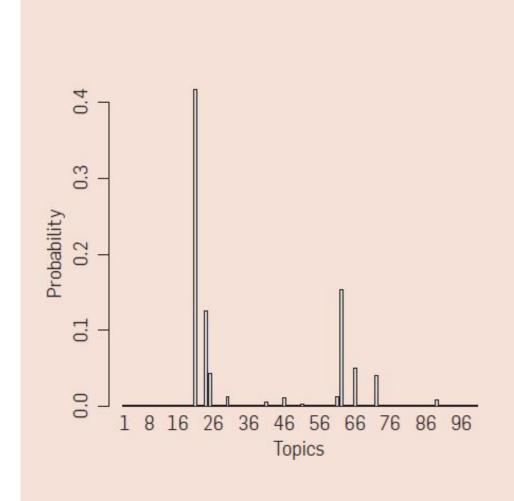
- For example, let's say we have three topics: sports, business, and science.
- For each topic, we uncover the most likely words to come from them:

For each word and topic pair, we learn some probability:
P(word|topic).

- The *topic distribution* is a multinomial distribution for each document representing what topics are most likely to appear in that document.
- For all our sample of documents, we have a distribution over {sports, science, business}.

```
ESPN article: [sports: 0.8, business: 0.2, science: 0.0]
Bloomberg article: [business: 0.7, science: 0.2, sports: 0.1]
```

For each topic and document pair, we learn some probability, P(topic | document).



"Genetics"	"Evolution"	4
human	evolution	
genome	evolutionary	
dna	species	
genetic	organisms	
genes	life	ı
sequence	origin	
gene	biology	
molecular	groups	
sequencing	phylogenetic	
map	living	
information	diversity	
genetics	group	
mapping	new	
project	two	
sequences	common	tı

"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

- Topic models are useful for organizing a collection of documents and uncovering the main underlying concepts.
- There are many variants that attempt to add even more structure to the 'model':
 - a. Supervised topic models guide the process with pre-decided topics.
 - b. Position-dependent topic models ignore which words occur in which document and instead focus on *where* they occur.
 - c. Variable number topic models test different numbers of topics to find the best model.

ACTIVITY: KNOWLEDGE CHECK

ANSWER THE FOLLOWING QUESTIONS



- 1. Take any recent news article and brainstorm which three topics this story is most likely to be made up of.
- 2. Next, brainstorm which words are most likely derived from which of those three topics.

DELIVERABLE

Topics and word-topic pairs

- gensim is a library of language processing tools focused on latent variable models of text.
- It was originally developed by grad students dissatisfied with current implementations of latent models.
- Documentation and tutorials are available on the <u>package's website</u>.

Let's first translate a set of documents (articles) into a matrix representation with a row per document and a column per feature (word or n-gram).

```
from sklearn.feature_extraction.text import CountVectorizer
cv = CountVectorizer(binary=False,
                     stop_words='english',
                     min df=3)
docs = cv.fit_transform(data.body.dropna())
# Build a mapping of numerical ID to word
id2word = dict(enumerate(cv.get_feature_names()))
```

- We want to learn which columns are correlated (i.e. likely to come from the same topic).
- This is the word distribution.
- We can also determine what topics are in each document, the *topic* distribution.

```
from gensim.models.ldamodel import LdaModel
from gensim.matutils import Sparse2Corpus

# First we convert our word-matrix into gensim's format
corpus = Sparse2Corpus(docs, documents_columns = False)

# Then we fit an LDA model
lda model = LdaModel(corpus=corpus, id2word=id2word, num topics=15)
```

- In this model, we need to explicitly specify the number of topic we want the model to uncover.
- This is a critical parameter, but there isn't much guidance on how to choose it. Try to use domain expertise where possible.

- Now we need to assess the *goodness of fit* for our model.
- Like other unsupervised learning techniques, our validation techniques are mostly about interpretation.
- Use the following questions to guide you:
 - Did we learn reasonable topics?
 - Do the words that make up a topic make sense?
 - Is this topic helpful towards our goal?

- We can evaluate fit by viewing the top words in each topic.
- gensim has a show_topics function for this:

```
num topics = 25
num words per topic = 5
for ti, topic in enumerate(lda.show topics(num topics = num topics,
num words per topic = n words per topic)):
    print("Topic: %d" % (ti))
    print (topic)
    print()
```

Some topics will be clearer than others. The following topics represent clear concepts:

```
0.009*cup + 0.009*recipe + 0.007*make + 0.007*food + 0.006*sugar \rightarrow Cooking and Recipes
```

```
0.013*butter + 0.010*baking + 0.010*dough + 0.009*cup + 0.009*sugar \rightarrow Cooking and recipes
```

0.013*fashion + 0.006*like + 0.006*dress + 0.005*style \rightarrow Fashion and Style

ACTIVITY: KNOWLEDGE CHECK

ANSWER THE FOLLOWING QUESTIONS



- 1. Demonstrate the code you used to generate the topics above.
- 2. Hypothesize other topic interpretations.

DELIVERABLE

Code and topics