

Feature LDA: a Supervised Topic Model for Automatic Detection of Web API Documentations from the Web

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- Introduction
- The Feature LDA model
- Data
- Experimental Results
- Conclusion



- What are Web APIs?
 - New Web Services based on a simple stack of technologies, ≈ "URL+HTTP+XML/JSON"
 - Known as RESTful service when conforming the REST principles
- Why Web APIs?
 - Light technology stack VS. "classical" Web services (WSDL, SOAP, WS-*)
 - Enable easy access and aggregation of collection of resources
 - Widely used and reused

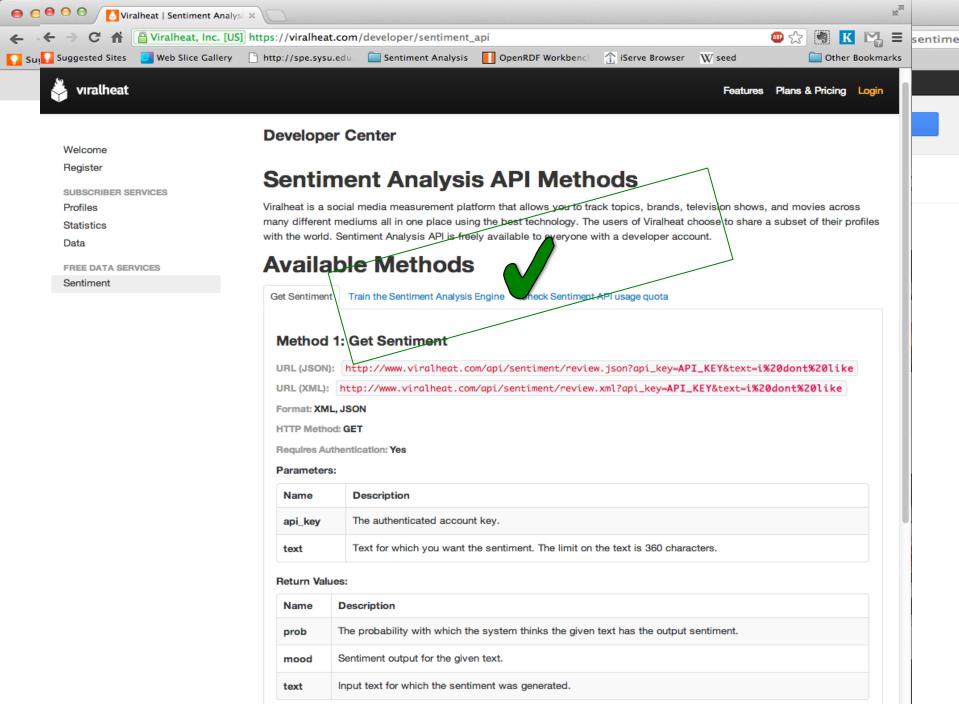


Finding a Web API

- Dedicated registries, e.g. ProgrammableWeb
 - Contain out of date or incorrect information, e.g. invalid pages or incorrect links to APIs documentation pages
 - Only a limited number of Web APIs listed, left out a large number of third party Web APIs
- General search engine, e.g. Google

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- Not optimized for Web API discovery
- Mix up with pages that are not (so) relevant,
 e.g. blogs and advertisement about Web APIs.



(http://www.ciarabnoge.com/ 2, Lympix (http://www.iymbix.com/ 2), etc.

www.guora.com/What-are-the-new-commercial-applications-for-text-analytics



- **Goal**: To build a customized search engine for detecting third party Web APIs on the Web scale
 - Assume every Web API provides public documentation page(s)
 - These pages provide the most relevant information for developers
 - Approached as a binary classification problem, i.e. distinguishing <u>API documentation</u> VS. <u>normal pages</u>



• Issues

- No simple way to effectively and uniquely identify Web APIs
- described in plain and unstructured HTML highly heterogeneous in format and contents, i.e. NO <u>Gold standard</u>
 - People hardly follow even there is !!!
- More than 99% of pages on the Web are NOT relevant to Web API
 - Need a high precision classifier yet maintaining good accuracy



The Feature LDA model

Latent Dirichlet Allocation (LDA)

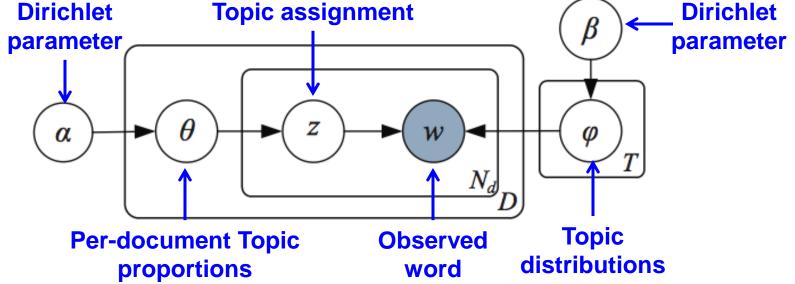
- LDA: the simplest form of topic models
 - A fully unsupervised Bayesian model
 - Assumes that documents exhibit multiple to known as "theme" or "gist")
 - Each topic is a distribution over words which tight semantic relation with one another

gene dna cell sequence genetics mapping human molecular

. . .

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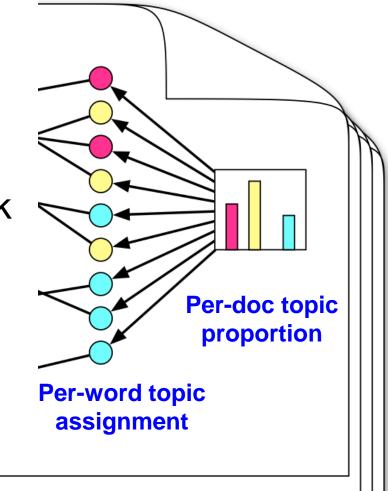
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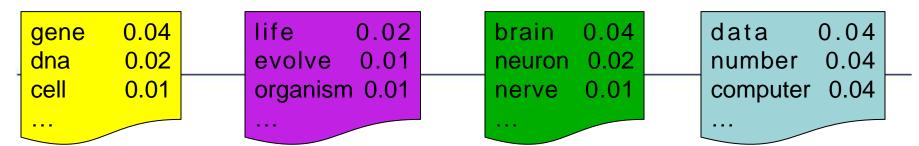
- Intuition:
 - Each document exhibit multiple topics
 - Each topic is a distribution over words
 - Each word is drawn from one of those topics

????...

Generate a document with a bulk of words ...



Topics:



Seeking Life's Bare (Genetic) Necessities

Haemophilus

genome 1703 genes

> Genes in common 233 genes

> > tycoplasma

genome

469 genes

COLD SPRING HARBOR, NEW YORK— How many genes does an organism peed 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 humangenome, notes Siv Andersson of Oppsala 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

parasite-specific

Minimal

gene set

250 genes

genes remov

Related and

-122 genes

128

genes

Ancest

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

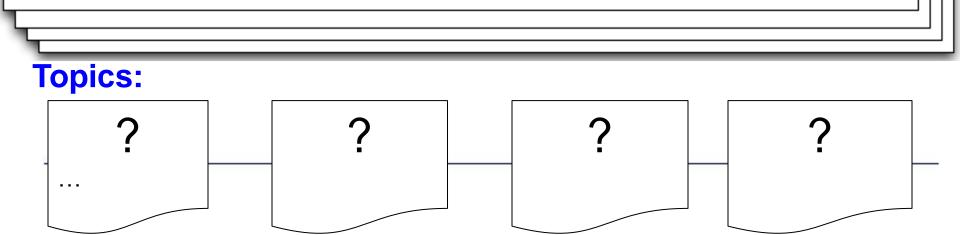
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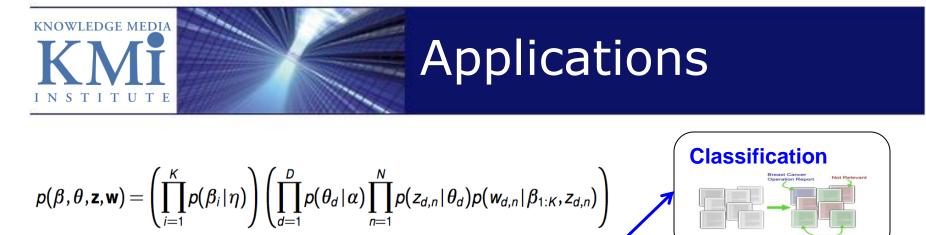
genes

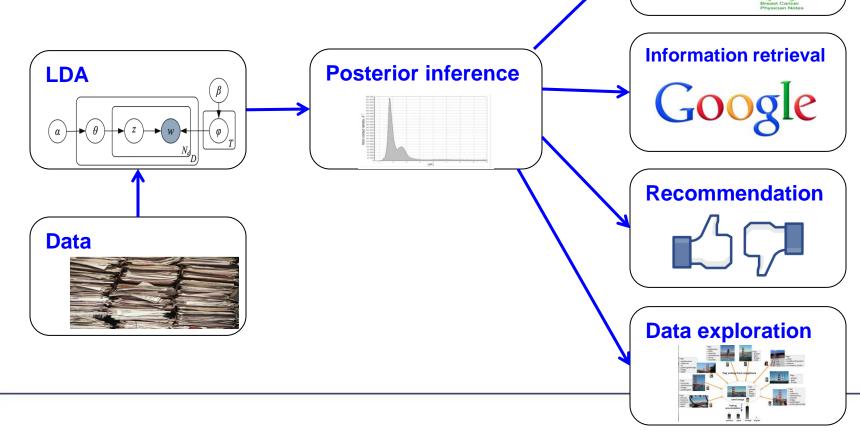
Genes needed

for biochemical pathways

SCIENCE • VOL. 272 • 24 MAY 1996





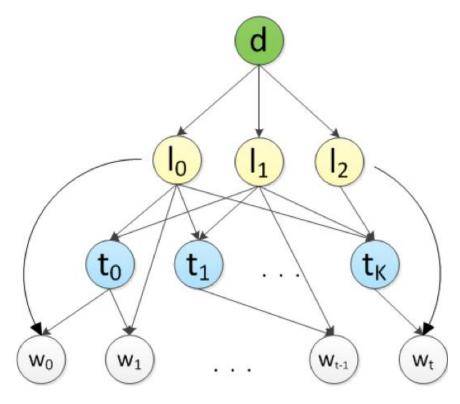


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- Feature LDA model: a generic probabilistic framework for text classification.
 - A <u>supervised</u> four-layer hierarchical Bayesian model
 - Accommodate supervisions from both <u>labelled</u> <u>instance</u> and <u>labelled features</u> for training
 - Able to extract meaningful class specific topics
- Labelled features
 - In *baseball* vs. *hockey* text classification
 - pitcher \rightarrow baseball, puck \rightarrow hockey
 - learned automatically from training data using any feature selection method, e.g. Info Gain

Feature LDA KNOWLEDGE MEDIA Graphical Model NSTITUTE Labelled **Topic** Per-doc class label features assignment Dirichlet specific topic **Observed** parameter proportions word λ Dirichlet parameter ß α φ ZK π **Document** N_d ε **Class label** class labels D specific topic **Per-doc class Class label** distributions proportions assignment

Generative Process



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For each document d

- Draw $\pi_d \sim Dir(\gamma \times \varepsilon_d)$
- For each class label k, draw $\theta_{d,k} \sim Dir(\alpha_k)$

For each word w in d

- Draw a class label $l_i \sim Mult(\pi_d)$
- Draw a topic $z_i \sim Mult(\Theta_{d,li})$
- Draw a word $w_i \sim \text{Mult}(\Phi_{li,zi})$



Collapse Gibbs sampling for model posterior estimation

$$P(z_t = j, c_t = k | \mathbf{w}, \mathbf{z}^{-t}, \mathbf{c}^{-t}, \boldsymbol{\alpha}, \boldsymbol{\beta}, \boldsymbol{\gamma}) \propto \frac{N_{k,j,w_t}^{-t} + \beta_{k,j,t}}{N_{k,j}^{-t} + \sum_i \beta_{k,j,i}} \cdot \frac{N_{d,k,j}^{-t} + \alpha_{k,j}}{N_{d,k}^{-t} + \sum_j \alpha_{k,j}} \cdot \frac{N_{d,k}^{-t} + \gamma_k}{N_d^{-t} + \sum_k \gamma_k}$$

• Approximating model parameters

$$\varphi_{k,j,i} = \frac{N_{k,j,i} + \beta_{k,j,i}}{N_{k,j} + \sum_i \beta_{k,j,i}} \quad \theta_{d,k,j} = \frac{N_{d,k,j} + \alpha_{k,j}}{N_{d,k} + \sum_j \alpha_{k,j}} \quad \pi_{d,k} = \frac{N_{d,k} + \gamma_k}{N_d + \sum_k \gamma_k}$$

Data and Setup

- The API dataset: 1,547 Web pages crawled from the <u>API Home URLs</u> of ProgrammableWeb (manually labelled, training/testing split: 80%-20%)
 - **622** pages are API documentations
 - 925 pages are normal Web pages
- Preprocessing
 - Extract content from HTMLs by discarding tags and java scripts that are not relevant to classification
 - remove wildcards, non-alphanumeric characters and stop-words, followed by Porter stemming.
- Setup

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- Class label k = 2
- Topic number T=1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 20
- 29,000 labelled features (Info Gain)

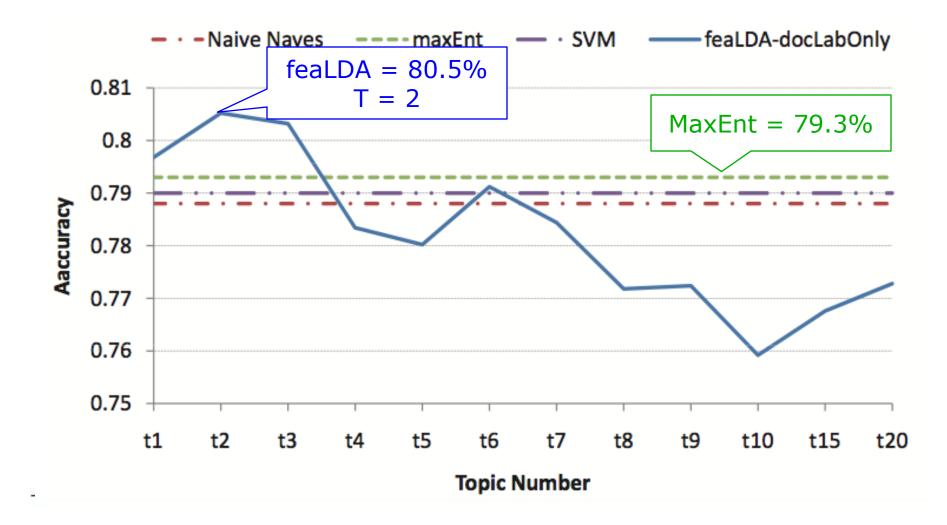




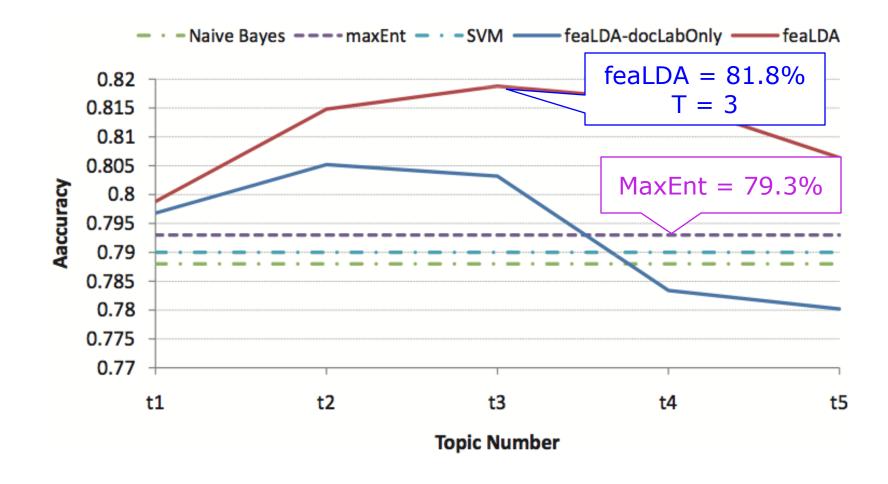
Experimental Results

- We report feaLDA classification results on the API dataset with different model settings:
 - Training with labelled instances
 - Training with both labelled instances and labelled features
 - feaLDA performance feature selection on labeled features
 - feaLDA vs. baselines (NB, SVM, MaxEnt) and other supervised topic models (labelLDA, pLDA)
 - Topic extraction

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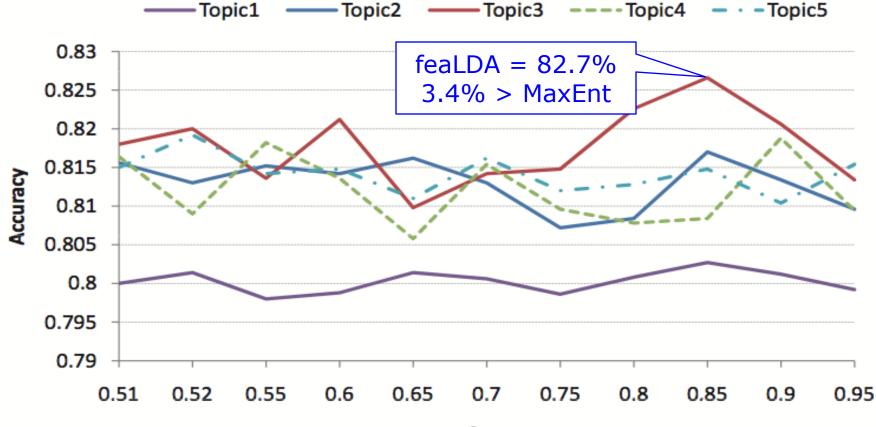
Labelled instances + labelled features



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 τ value

_ feaLDA classification accuracy vs. different feature class probability threshold τ.

Overall comparison

Table 2: Comparing feaLDA with existing supervised approaches.

	Naive Bayes	SVM 1	maxEnt	labeled LDA	pLDA	feaLDA
Recall	79.2	70.8	69.3	59.8	65.9	68.8
Precision	71.0	75.4	77.4	85.1	82.1	85.2
$\mathbf{F1}$	74.8	73.1	73	70.2	73.1	76
Accuracy	78.6	79	79.3	79.8	80.5	82.7

- feaLDA vs. the state-of-the-art models
 - outperforms three strong supervised baselines
 - better than labeledLDA and pLDA for more than 3% in accuracy
 - gives very high precision: essential for reducing false positives when mining from the Web

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Topic Extraction

Table 3: Topics extracted by featDA with K = 2, T = 3.

T1: nbsp quot gt lt http api amp type code format valu json statu paramet element
 T2: lt gt id type http px com true url xml integ string fond color titl date
 T3: api http user get request url return string id data servic kei list page paramet
 T1: px color font background pad margin left imag size border width height text div thread
 T2: servic api site develop data web user applic http get amp email contact support custom
 T3: obj park flight min type citi air fizbber airlin stream school die content airport garag

• Topics with true API label

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- Terms are fairly technical, e.g. json, statu, paramet, element, valu, request and string, etc.
- Topics with false API label
 - Terms are less technical and more diverse, e.g. contact, support, custom, flight, school



- Discovering Web APIs is becoming increasingly important and existing support is not optimal
- Treat Web API discovery as a classification problem
- Presented a supervised topic model called feaLDA
 - offers a generic framework for text classification
 - Capable to encode supervision from both labelled instance and labelled features
 - Offers very high precision which is crucial for reducing false positive when mining from the Web
 - Able to extract class label specific topics



Questions?

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