



Classification tasks

Clustering

 classifying the given data into k clusters by defining k centroids
 minimizing a chosen Euclidean distance between a data point

and cluster center

(PDF) BamBam: Genome sequence analysis tools for biologists



- 1. Random choice
- 2. Distance from each data point to the centroid
- 3. Closer ones belong to the same cluster
- 4. New centroid of each cluster (average)
- 5. Repeat 2-4
- 6. Till the the centroid value stay the same











Cluster Number

Perplexity 1.

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2. Semantic Coherence Metric





Cluster Number 1. Perplexity

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2. Semantic Coherence Metric



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Topic Modeling

Document Representation

Vector Space Model:

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- Term frequency (in a doc)
- Term Frequency–Inverse Document Frequency (TF-IDF)

Document Length

Long:

- Latent Semantic Analysis (LSA): doc-topic & topic-term matrix
- Latent Dirichlet Allocation (LDA):

Gibbs sampling: the distance between words in the same latent topic is minimized & the distance between words from different latent topics is maximized. · • •

Topic Modeling

Document Length

Challenges

- Data sparsity
- Context
- Labeled data
- Noises 🦥 🦦 🦭 🐋

Short:

- Sentence-LDA: Each document is inferred from only one topic
- Global Word Co-Occurrences
 Based Methods:
 - Documents with similar context tend to share the same topics
- FastText-based Sentence-LDA: FastText associates each word with a group of similar words with a similarity degree or weight.

The main hypothesis of our proposed model is that a document can be about several topics.

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Case StudyUsing Topic Modeling and
Word Embedding for Topic
Extraction in Twitter -
ScienceDirectLDA+W2V:Image: Image: I

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For the data we used 114826 tweets as our short text documents and to collect them we used Twitter API¹. To label the topics of the collected data we used a semi-automatic annotation technique. The main idea behind the latter is to try to cluster documents or tweets, using k-means, in a way documents within the same cluster are similar to each other. After clustering our documents, we will try to take a few documents (sample) from each cluster and annotate them manually. After the manual annotation, all documents within the same cluster from which we took the sample will take the label of its correspondent sample since all documents in one cluster share similar properties (see figure3).



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ScienceDirectLDA+W2V:Image: Case StudyImage: Case StudyTopic-doc matrixImage: Case StudyImage: Case Study

After calculating the document's score in the range of the detected topics (topics detected in section 3.4.1) we obtain as a result a vector where each row represents a topic, and the column-row values represent the score of the document in the row's topic. From a topics' vector of document *d*, we will extract the most representative topics by fixing a threshold *th*. So, if the topic's score is greater than *th* so the latter is one of document *d*'s relevant topics.

 $relevant_topics(d, th) = \{t \in Topics : withdoc_topic_score(d, t) \ge th\}$

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A good threshold value :

- maximizes the percentage of good predictions
- minimizes the number of predictions labels per tweet

(2)

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The best value to choose is 0.5

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- minimizes the number of labels "3"
- maximizes the threshold values (topics are relevant)



(2)

