# *litsift*: Automated Text Categorization in Bibliographic Search

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## 1 Motivation

- **Goal:** comparison of computational results from bioinformatics with experimental results from life sciences
- **Task:** find relevant literature containing information on *conserved RNA secondary structures in viral genomes* for a fixed virus group

#### **Complications:**

- relevant results may be hidden in articles with differing main topics
- key words may be omitted because context is clear or may be overloaded (e.g. secondary structure)
- no established nomenclature of RNA features in viruses
- ⇒ Exploratory Project: assess the feasibility of supporting broad bibliographic search with automated text categorization techniques (2PM).

# 2 Approach

- 1. learn relevant literature using training corpus (dedicated to a specific virus group, e.g. *Picornaviridae, Flaviviridae*)
- 2. create test corpus (on some other virus group) by searching bibliographic database and downloading referenced articles
- 3. apply trained classifier to test corpus
- 4. present articles as ranked list
- 5. manually relabel some test articles and use for retraining





#### **3 Data Sets**

Corpus	Source	Size	Positive
picorna	Pubmed	40	68%
picorna2	Pubmed + Experts	64	58%
flavi	Pubmed	153	8%
flavi2	Pubmed + Experts	187	12%
hepadna	Pubmed	16	69%

## 4 Methods

#### 4.1 Data Preparation

- 1. download: Perl wrapper scripts
- 2. PDF  $\rightarrow$  Text conversion: pdftotext, ps2ascii
- 3. tokenization and full text index: ConceptComposer
- 4. term relevance measures: SQL script

• Odds Ratio 
$$OR(t,c) = \frac{P(t|c) \cdot (1 - P(t|\overline{c}))}{(1 - P(t|c)) \cdot P(t|\overline{c})}$$
  
• Mutual Information  $MI(t,c) = \log \frac{P(t,c)}{P(t,c)}$ 

• Mutual Information 
$$MI(t,c) = \log \frac{I(t,c)}{P(t) \cdot P(c)}$$

5. vector representation: SQL script, using tfidf term weights (persistent storage: MySQL relational database)

#### 4.2 Automated Text Categorization

**Prototype:** Java application on top of Weka 3 and MySQL. Supports crossvalidation on training corpus and validation on separate test corpus. External data download, preparation, and labeling.

Parameters for experiments:

- term relevance measure:  $\{OR, MI\}$
- dimensionality:  $\{10, 20, ..., 200\}$
- target recall: {80%}
- classifier type {SMO, J48, N.B.} (i.e., SVM, C4.5, Naive Bayes)
- classifier-specific parameters

## **5** Results

#### **5.1 Feature Selection**

Relevance measure used as classifier. Threshold defined by target recall 100%. Average precision:

$p_{avg}$	flavi	flavi2	picorna	picorna2
OR	7.8%	11.8%	67.6%	58.0%
MI	11.2%	20.2%	76.7%	69.3%

 $\Rightarrow$  baseline for cross evaluation.

#### 5.2 Cross Evaluation

**Picorna** corpora: easy to classify. E.g., SMO with MI on picorna2:



Flavi corpora: harder to classify. E.g., SMO with MI on flavi2:



Cross Validation Performance (Experiment #105)

Typically less than 50 features needed for maximum precision.

#### 5.3 Validation on Separate Test Corpus

Classifiers trained on Flavi corpora transfer well to Picorna corpora (e.g., SMO with OR, flavi2  $\rightarrow$  picorna2)...

1 0.8 precision / recall 0.6 0.4 0.2 Precision Recall 0 20 40 120 160 180 200 60 80 100 140 0 # features

Validation Performance (Experiment #66)

... but not vice versa (e.g., SMO with OR, picorna2  $\rightarrow$  flavi2)



Still, even a low precision may save work...

## 6 Cost Model

- **Task:** find at least a fraction r of all relevant documents within a bibliographic search result, i.e., target recall is r.
- **Goal:** minimize fraction q of articles to be inspected manually.
- **Baseline:** random selection with probability r requires  $q_{rand} = r$  and yields recall r.

With classifier: classifier with precision p requires  $q_{auto} = \min(P(c)r/p, 1)$  where P(c) frequency of relevant documents

Work reduction:  $s = (q_{rand} - q_{auto})/q_{rand} = 1 - P(c)/p$  if  $P(c) \le p$ 

#### 6.1 Work Reduction (Examples)

training	test	class	msr	P(c)	pmax	r	S
flavi2	picorna2	SMO	MI	58%	83.3%	100.0%	30%
picorna2	flavi2	SMO	OR	12%	32.7%	81.8%	63%
flavi2	hepadna	SMO	OR	69%	90.9%	90.9%	25%
picorna2	hepadna	SMO	OR	69%	90.0%	81.8%	25%

## 7 Conclusions

- classifiers can be transferred among corpora on different virus groups, at the cost of reduced precision
- low precision can still reduce manual work significantly, especially with infrequent classes
- work reduction allows to broaden search queries and to increase overall recall

#### 8 Future Plans

- experiment with classifiers for partially unlabeled data sets
- complete implementation of *litsift* tool:
  - implement Web interface based on Apache Cocoon
  - re-implement download manager in Java, based on Apache xalan and JaxME.