Evaluating Feature Selection Methods for Multi-Label Text Classification

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Motivation

• Real word, exciting research problem on large-scale biomedical semantic indexing



• Can feature selection help?

Multi-Label Learning

• Multi-label data

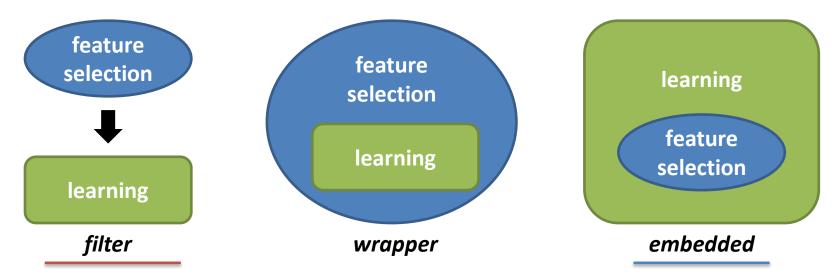
- Instances related with a subset of a finite label set

| | Pain | Fever | ••• | Weight | Disease |
|-----------|------|-------|-----|--------|--------------------------------------|
| Patient 1 | yes | no | | 101.5 | {gastritis, duodenitis} |
| Patient 2 | no | yes | | 61.2 | {esophagitis} |
| : | : | : | •. | : | : |
| Patient M | yes | yes | ••• | 79.8 | {esophagitis, gastritis, duodenitis} |

- Models learned from such data can output
 - Bipartition of label set, ranking of labels, ranking of instances, marginal/joined probabilities

Feature Selection

- Main objectives
 - Reducing measurement & storage requirements, data understanding, reducing training & utilization times, improving prediction accuracy
- Three main categories of approaches



Multi-Label Filter Feature Selection

- Step 1: Feature ranking separately per label
 One can use any standard single-label feature evaluation measure for binary classification
- Step 2: Aggregation of the different rankings
 - Mean, Max of the evaluation score for all labels
 - Round Robin (RoR), Rand Robin (RaR) selection
 per label based on the evaluation scores

Example – Mean Aggregation

| Feature | Score <i>Y</i> ₁ | Score Y ₂ | Score <i>Y</i> ₃ | Mean | Ranking |
|-----------------------|-----------------------------|----------------------|-----------------------------|------|-----------------------|
| <i>X</i> ₁ | 0.1 | 0.9 | 0.5 | 0.5 | X_5 |
| <i>X</i> ₂ | 0.6 | 0 | 0.3 | 0.3 | <i>X</i> ₃ |
| <i>X</i> ₃ | 0.5 | 0.7 | 0.6 | 0.6 | <i>X</i> ₁ |
| X_4 | 0.3 | 0.5 | 0.4 | 0.4 | X_4 |
| <i>X</i> ₅ | 0.7 | 0.6 | 0.8 | 0.7 | <i>X</i> ₂ |

Example – Max Aggregation

| Feature | Score <i>Y</i> ₁ | Score Y ₂ | Score <i>Y</i> ₃ | | Max | Ranking | Mean | |
|-----------------------|-----------------------------|----------------------|-----------------------------|--|-----|---------|-----------------------|-----------------------|
| <i>X</i> ₁ | 0.1 | 0.9 | 0.5 | | | 0.9 | X_1 | X_5 |
| <i>X</i> ₂ | 0.6 | 0 | 0.3 | | 0.6 | X_5 | <i>X</i> ₃ | |
| <i>X</i> ₃ | 0.5 | 0.7 | 0.6 | | | 0.7 | <i>X</i> ₃ | <i>X</i> ₁ |
| X_4 | 0.3 | 0.5 | 0.4 | | | 0.5 | <i>X</i> ₂ | X_4 |
| <i>X</i> ₅ | 0.7 | 0.6 | 0.8 | | 0.8 | X_4 | <i>X</i> ₂ | |

Example – RoR Aggregation

| | $\downarrow\downarrow$ | $\downarrow\downarrow\downarrow$ | \downarrow |
|-----------------------|-----------------------------|----------------------------------|-----------------------------|
| Feature | Score <i>Y</i> ₁ | Score Y ₂ | Score <i>Y</i> ₃ |
| <i>X</i> ₁ | 0.1 | 0.9 | 0.5 |
| <i>X</i> ₂ | 0.6 | 0 | 0.3 |
| <i>X</i> ₃ | 0.5 | 0.7 | 0.6 |
| X_4 | 0.3 | 0.5 | 0.4 |
| X_5 | 0.7 | 0.6 | 0.8 |

| Ranking |
|-----------------------|
| X_5 |
| <i>X</i> ₁ |
| <i>X</i> ₂ |
| <i>X</i> ₃ |
| X_4 |

Example – RaR Aggregation

| | $\downarrow\downarrow\downarrow$ | \downarrow | $\downarrow\downarrow\downarrow$ |
|-----------------------|----------------------------------|----------------------|----------------------------------|
| Feature | Score <i>Y</i> ₁ | Score Y ₂ | Score <i>Y</i> ₃ |
| <i>X</i> ₁ | 0.1 | 0.9 | 0.5 |
| <i>X</i> ₂ | 0.6 | 0 | 0.3 |
| <i>X</i> ₃ | 0.5 | 0.7 | 0.6 |
| X_4 | 0.3 | 0.5 | 0.4 |
| <i>X</i> ₅ | 0.7 | 0.6 | 0.8 |
| frequency | 0.3 | 0.1 | 0.4 |

| Ranking | RoR |
|-----------------------|-------|
| <i>X</i> ₅ | X_5 |
| <i>X</i> ₃ | X_1 |
| <i>X</i> ₁ | X_2 |
| <i>X</i> ₂ | X_3 |
| X_4 | X_4 |

Experimental Setup (1/2)

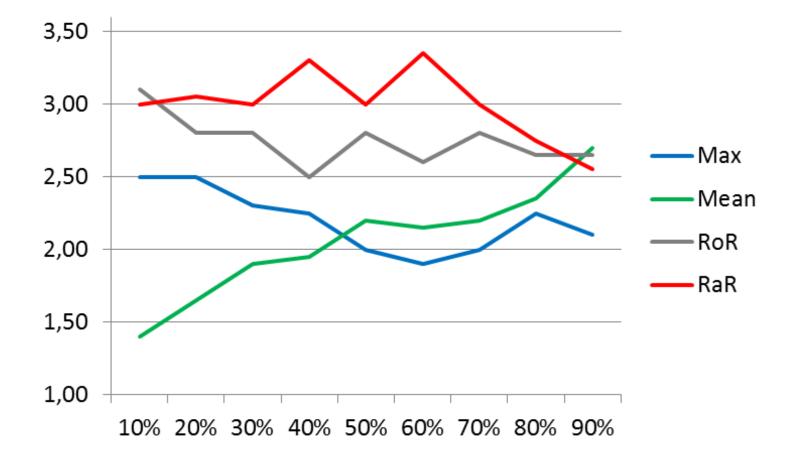
- 20 benchmark textual datasets
 - yahoo (11), enron, delicious, bookmarks, bibtex, medical, tmc007, slashdot, language log, rcv1v2
- 8 filter feature selection methods
 - 2 feature evaluation measures (χ^2 , BNS)
 - 4 aggregation strategies (Mean, Max, RoR, RaR)
- 2 baselines

- random feature selection (RFS), all features (AF)

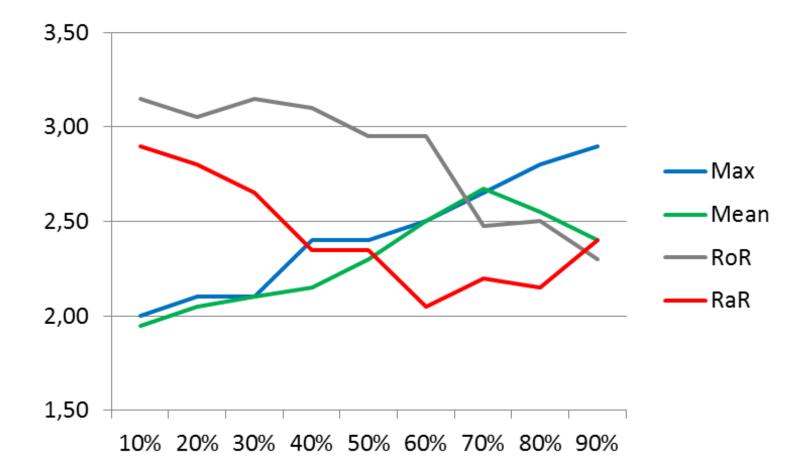
Experimental Setup (2/2)

- Multi-label classification
 - Binary Relevance (*aka one-vs-rest*) with linear support vector machines as base algorithm
- Evaluation
 - Micro F-measure
 - Selection of 10%, 20%, ..., 90% features
 - Average ranking of methods across datasets

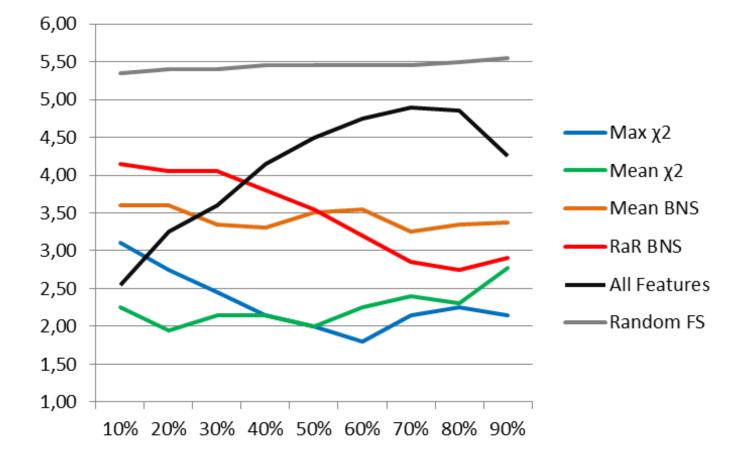




BNS Results



Best of χ^2 and BNS, Random, All



Recap

- Empirical study with large number of text datasets (20) in contrast with past literature
- Aggregations RaR and RoR tried for the first time here, but did not work successfully
- BNS is worse than χ^2 , contrary to findings for single-label data
- For χ^2 mean (max) aggregation should be preferred for low (high) percentage of features

Future Work

- Binary relevance + global feature selection
- Binary relevance + local feature selection
- Meta-labeler + global feature selection

 First results on BioASQ data are negative
 Will verify this on the 20 datasets of this study
- Meta-labeler + local feature selection
 Fails, as it renders the SVM scores incomparable
- Explore efficient ways to exploit label dependence in multi-label feature selection

The End

- Thank you for your attention!
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