

# Outline

- Intro: Multi-label Classification
- RM: The *R*anking *M*ethod
- A Probability Distribution for RM
- END: *E*nsembles of *N*ested *D*ichotomies
- Related multi-label work: Pairwise Training
- Current work: M-END: *M*ulti-label END
- Current work: EsPS: *E*nsembles of *s*plit *P*runed *S*ets
- Summary

# Intro: Multi-label Classification

*Single-label (Multi-class) Classification:*

- Set of instances  $D$ . Set of labels (classes)  $L$ .
- For each  $d \in D$ , select a label (class)  $l \in L$
- Single-label representation:  $(d, l)$

*Multi-label Classification:*

- Set of instances  $D$ . Set of labels  $L$ .
- For each  $d \in D$ , select a **label subset**  $S \subseteq L$
- **Multi-label** representation:  $(d, S)$

e.g.  $L = \{Sports, Environment, Science, Politics\}$

(“Revealed: Polluting impact of humans on the oceans...”,  $\{Environment, Science\}$ )

# Intro: Multi-label Classification

**Problem Transformation:** Multi-label problems are transformed into one or more single-label problems.

**Algorithm Adaption:** Employs Problem Transformation internally to a single-label algorithm.

i.e. All multi-label classification involves **Problem Transformation**. There are three fundamental methods:

- BM (Binary Method)
- RM (Ranking Method)
- CM (Combination Method)

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# RM: The Ranking Method

Using the probability distribution from a single-label classifier, and a threshold ( $t$ ), multi-labels are selected.

$L = \{Sports, Environment, Science, Politics\}$

ML $D_{train}; (d, S \subseteq L)$
$d_1, \{Sports, Politics\}$
$d_2, \{Science, Politics\}$
$d_3, \{Sports\}$
$d_4, \{Environment, Science\}$

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SL  $D_{train}; (d, l \in L)$

$d_1, Sports$

$d_1, Politics$

$d_2, Science$

$d_2, Politics$

$d_3, Sports$

$d_4, Science$

$d_4, Environment$

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$(l \in L)$	$P(l d_x)$
<i>Sports</i>	$\lambda$
<i>Environment</i>	$\lambda$
<i>Science</i>	$\lambda$
<i>Sports</i>	$\lambda$

$d_x \in D_{test}$

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$(l \in L)$	$P(l d_x)$
<i>Sports</i>	0.03
<i>Environment</i>	0.48
<i>Science</i>	0.45
<i>Politics</i>	0.04

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$d_x \in D_{test}$  ,  $t = 0.2$

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$d_x \in D_{test}$  ,  $t = 0.2$  :  $(d_x, \{Environment, Science\})$

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$d_x \in D_{test}$  ,  $t = 0.2$  :  $(d_x, \{Environment, Science\})$

- Assumes that all labels are independent
- Issues with threshold selection / classifier selection

# RM: SL Classifier Selection

Each PT method must be supplied a single-label (SL) classifier

```
weka.classifiers.multilabel.RM -t MEDC.arff
```

e.g. SMO:

```
-W weka.classifiers.functions.SMO
```

e.g. SMO – $M$  (probabilistic outputs):

```
-W weka.classifiers.functions.SMO -M
```

e.g. SMO under Bagging:

```
-W weka.classifiers.meta.Bagging --
```

```
-W weka.classifiers.functions.SMO
```

e.g. Ensembles of Nested Dichotomies:

```
-W weka.classifiers.meta.END --
```

```
-W weka.....meta.nestedDichotomies.ND --
```

```
-W weka.classifiers.functions.SMO
```

# RM: SL Classifier Selection

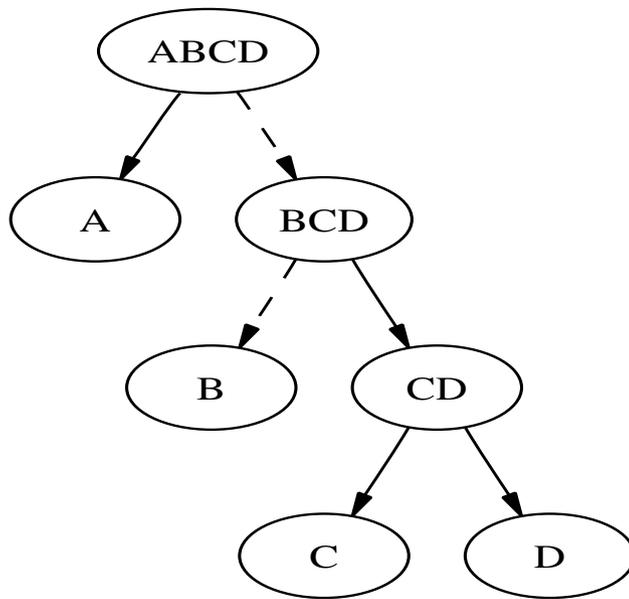
RM: Accuracy on the *Medical* and *Enron* datasets.

$D$	SMO	SMO $-M$	Bag. SMO	END		
				ND SMO	CBND SMO	DNBND SMO
MED	74.53	75.43	73.47	77.74	80.31	79.55
ENR	20.90	38.49	23.78	42.86	40.47	42.53

- *Medical*: average of 1.25 labels / instance
- *Enron*: average of 3.38 labels / instance
- In each case SMO is the SL classifier. Therefore the difference is in the probability distribution and the comparison between labels.

# END: Ensembles of Nested Dichotomies

- <sup>a</sup>: Single-label; Binary Tree; Dichotomy; Ensembles
- Can supply RM-transformed multi-label data and use prob. distribution and threshold to gather multi-labels



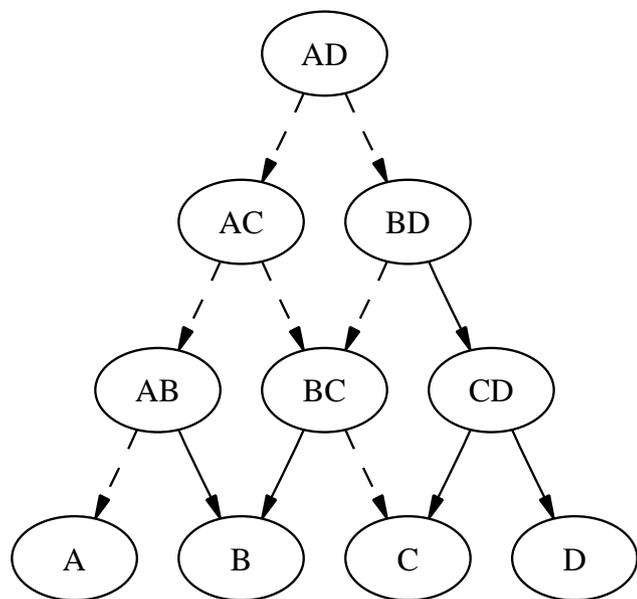
an example dichotomy,  $L = \{A, B, C, D\}$ .  $(d_x, B)$

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<sup>a</sup>Eibe F. and Kramer S. Ensembles of nested dichotomies for multi-class problems. Proc. of the 21st International conference of Machine Learning, 2004.

# Classification by Pairwise Training

- <sup>a</sup>: Multi-label; DAG; Trichotomy

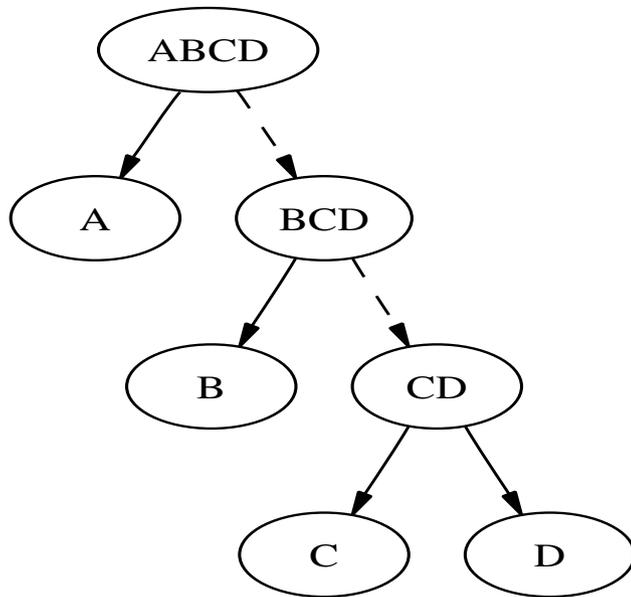


an example DAG,  $L = \{A, B, C, D\}$ .  $(d_x, \{A, C\})$

- Error propagation
- Classification sensitive to DAG arrangement (Ensembles?)

# M-END: Multi-label END

- Multi-label; Binary Tree; Dichotomy; Ensembles



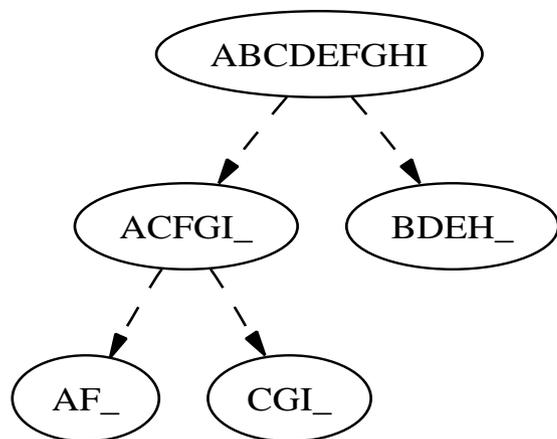
an example dichotomy,  $L = \{A, B, C, D\}$ .  $(d_x, C, D)$

- Can return multiple labels if these labels co-occur often
- Creates splits according to label co-occurrences

A work in progress...

# EsPS: *E*nsembles of *s*plit *P*runed *S*ets

- A somewhat related method inspired by work on M-END



$$L = \{A, \dots, I, \_ \} (\_ = null)$$

- Some param. to determine when branching stops
- Multi-label method (i.e. PS aka PPT) train on each leaf subset
- Classifications of all leaves are combined

A work in progress...

# EsPS: *E*nsembles of *s*plit *P*runed *S*ets

Compared to state-of-the-art RAKEL:

<i>Dataset</i>	RAKEL	CM	BM	RM	EsPS
Scene	71.58±0.89	71.81±1.22	58.28±0.92 ●	71.72±0.98	73.51±0.56 ○
Medical	72.55±2.32	74.71±1.32	73.00±1.08	72.71±1.56	76.86±1.60
Yeast	54.49±0.98	51.98±0.93 ●	49.64±0.88 ●	51.95±0.62 ●	54.21±0.95
Enron	42.98±0.63	41.02±1.08	38.64±1.05 ●	27.22±0.31 ●	47.55±0.80 ○
Reuters	31.80±0.29	49.17±0.67 ○	31.91±0.76	49.08±0.59 ○	46.52±1.03 ○

○, ● statistically significant improvement or degradation

<i>Dataset</i>	PS.END.DNBND
Medical*	75.82±1.52

5 x 2 fold CV; all ensemble methods 10 iterations, all other parameters tuned via 5 fold internal CV. SMO used as internal single-label classifier in each case.

# Summary

- Good probability distributions are important to threshold methods like RM
- Ensembles are a way to achieve this (plus the additional benefits of an ensemble)
  - END
- In multi-label classification it is very important to take into account label-correlations
  - M-END
  - *EsPS*