

# A Framework for Machine Learning with Ambiguous Objects

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The talk involves some joint work with my students :

*Min-Ling Zhang*

*Yin-Xing Li*

*Yu-Feng Li*

*Sheng-Jun Huang*

... ..

And my collaborators:

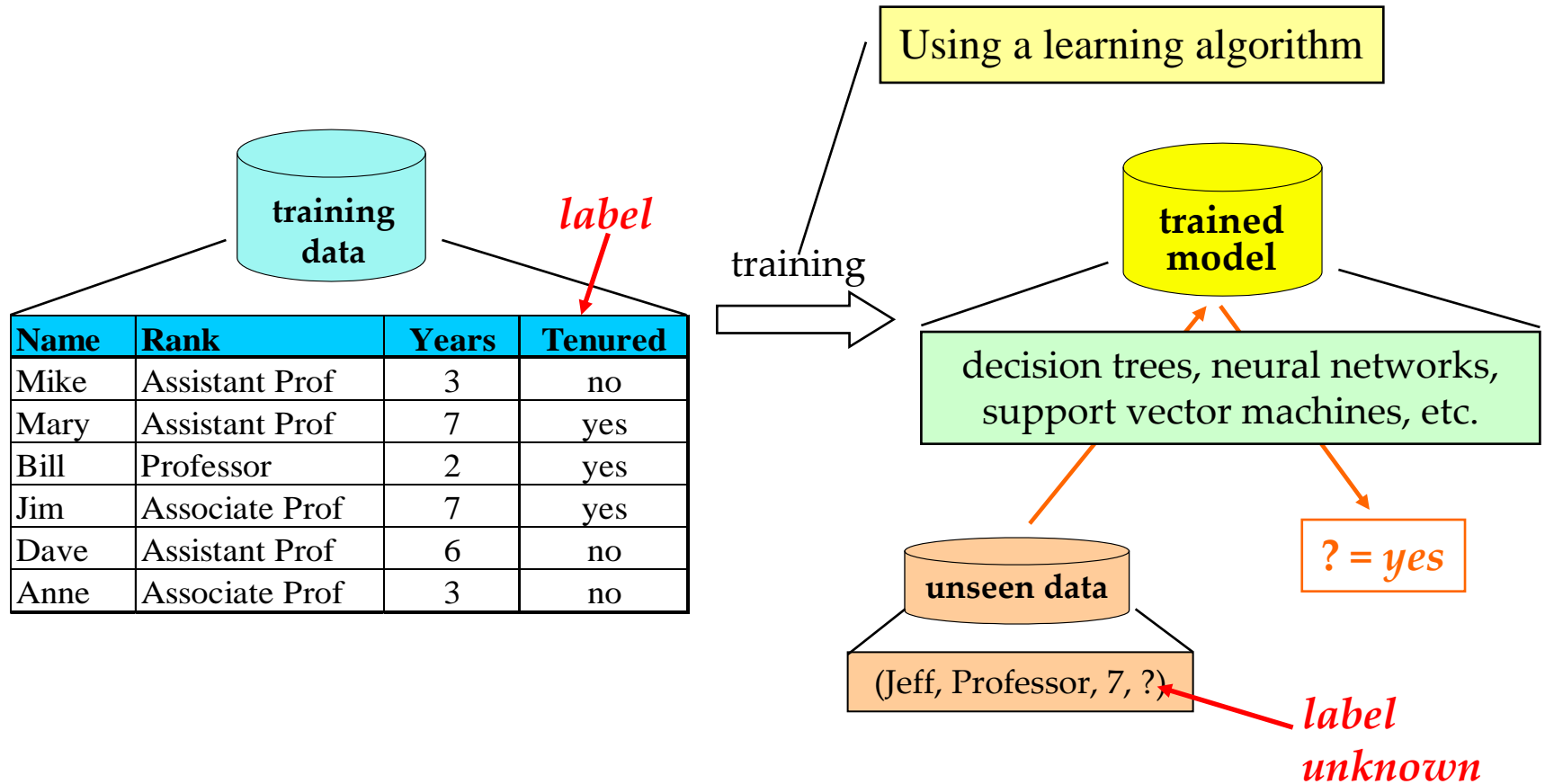
*Jieping Ye*

*Shuiwang Ji*

*Sudir Kumar*

... ..

# A typical machine learning process

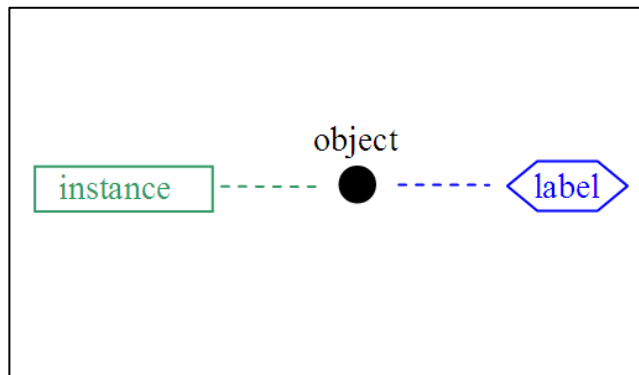


# Traditional Machine Learning Setting

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In traditional supervised learning:

- A real-world object is represented by an **instance** (feature vector)
- The instance is associated with a **label** which indicates the concerned characteristics (such as categorization) of the object



$\mathcal{X}$  - the instance space

$\mathcal{Y}$  - the set of class labels

## The task:

To learn a function  $f: \mathcal{X} \rightarrow \mathcal{Y}$  from a given data set  $\{(\mathbf{x}_1, y_1), (\mathbf{x}_2, y_2), \dots, (\mathbf{x}_m, y_m)\}$  where  $\mathbf{x}_i \in \mathcal{X}$  is an instance and  $y_i \in \mathcal{Y}$  is the known label of  $\mathbf{x}_i$

# Ambiguous Data

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*Elephant ?  
Tropic ?*

*Lion ?  
Africa ?*

*Grassland?  
... ..*

# Ambiguous Data

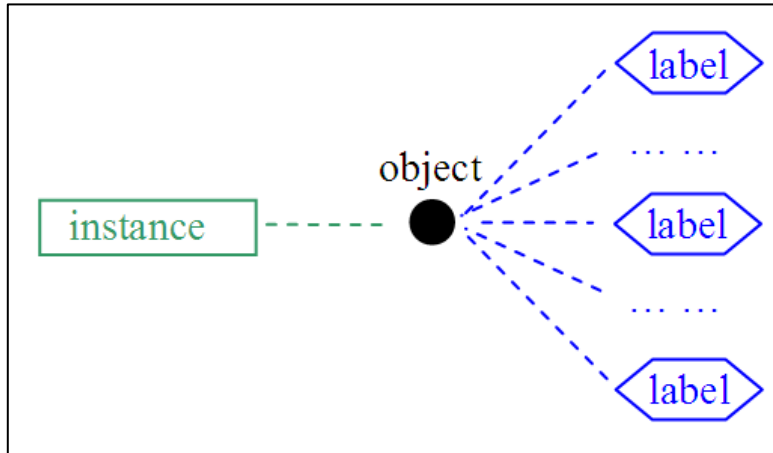
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*Scientific novel*  
*Jules Verne's writing*  
*Book on traveling*

... ..

# Multi-Label Learning



## MLL task:

To learn a function  $f_{MLL} : \mathcal{X} \rightarrow 2^{\mathcal{Y}}$  from a given data set  $\{(x_1, Y_1), (x_2, Y_2), \dots, (x_m, Y_m)\}$ , where  $x_i \in \mathcal{X}$  is an instance and  $Y_i \subseteq \mathcal{Y}$  is a set of labels  $\{y_1^{(i)}, y_2^{(i)}, \dots, y_{l_i}^{(i)}\}$ ,  $y_k^{(i)} \in \mathcal{Y}$  ( $k = 1, 2, \dots, l_i$ ).

$\mathcal{X}$  - the instance space

$\mathcal{Y}$  - the set of class labels

$l_i$  - the number of labels in  $Y_i$

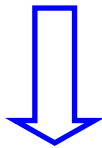
# Multi-Label Learning Algorithms

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- Decomposing the task into multiple binary classification problems each for a class
  - ✓ MLSVM [Boutell et al., PR04]
  - ✓ ... ..
  
- Considering the ranking among labels
  - ✓ BoosTexter [Schapire & Singer, MLJ00]
  - ✓ BP-MLL [Zhang & Zhou, TKDE06]
  - ✓ RankSVM [Elisseeff & Weston, NIPS'01]
  - ✓ ... ..
  
- Exploring the class correlation
  - ✓ Probabilistic generative models [McCallum, AAI'99w; Ueda & Saito, NIPS'02]
  - ✓ Maximum entropy methods [Ghamrawi & McCallum, CIKM'05; Zhu et al., SIGIR'05]
  - ✓ ... ..



# The Problem



$[x_1, x_2, \dots, x_d]^T$

**one-to-many  
mapping**

Elephant

Lion

Grassland

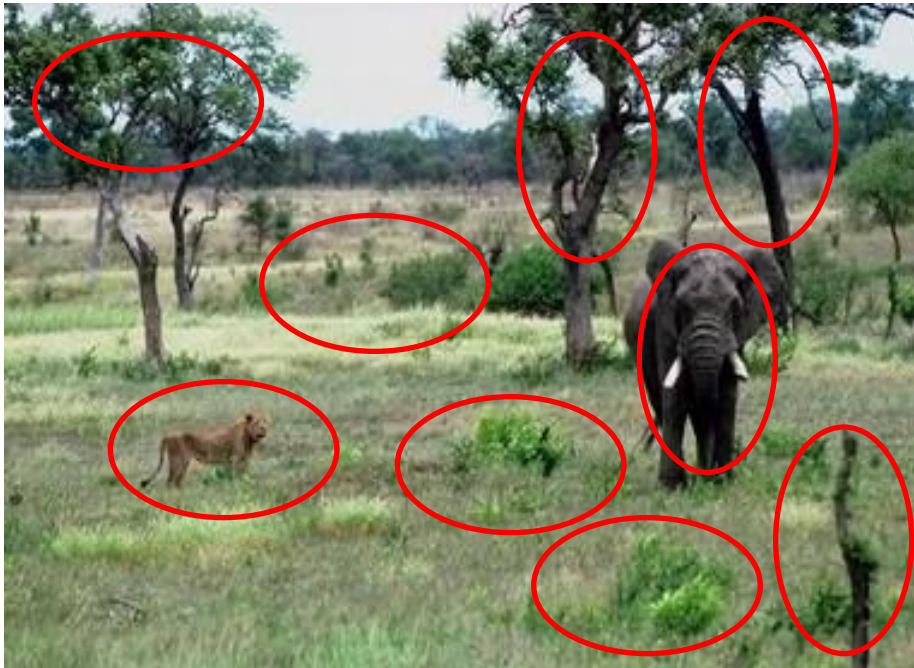
Tropic

Africa

## Consider ...

---

An image usually contains **multiple** regions each can be represented by an instance



The image can simultaneously belong to **multiple** classes

*Elephant*

*Lion*

*Grassland*

*Tropic*

*Africa*

... ..

## Consider ...

---

A document usually contains **multiple** sections each can be represented by an instance



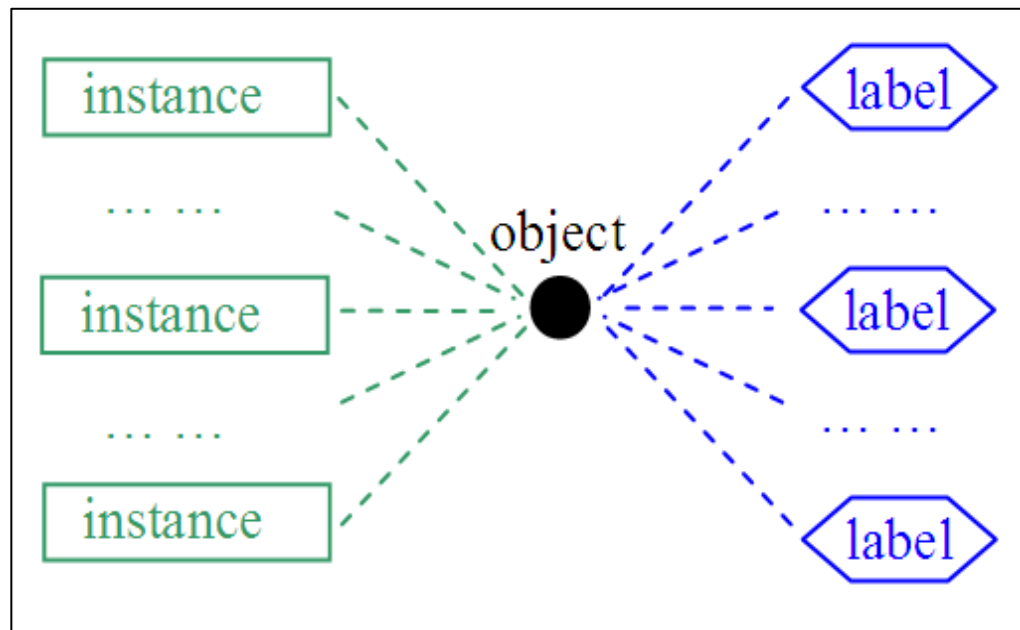
The document can simultaneously belong to **multiple** categories

*Scientific novel*

*Jules Verne's writing*

*Book on traveling*

... ..



## Multi-Instance Multi-Label (MIML) Learning 多示例多标记学习

# Why MIML ?

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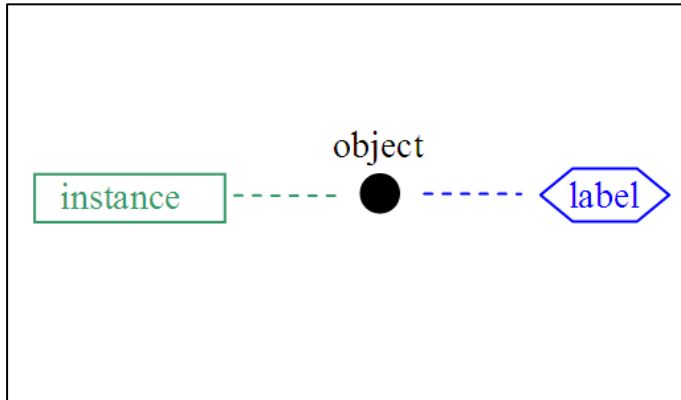
Appropriate representation is important

Having an appropriate representation is as important as having a strong learning algorithm

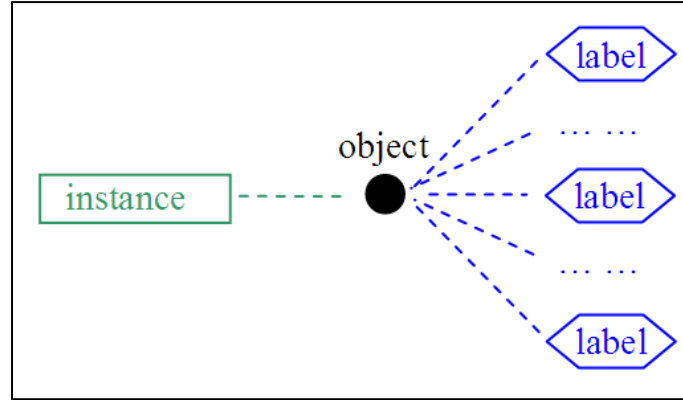
MIML captures more information of ambiguous data

Traditional supervised learning, multi-instance learning and multi-label learning are degenerated versions of MIML

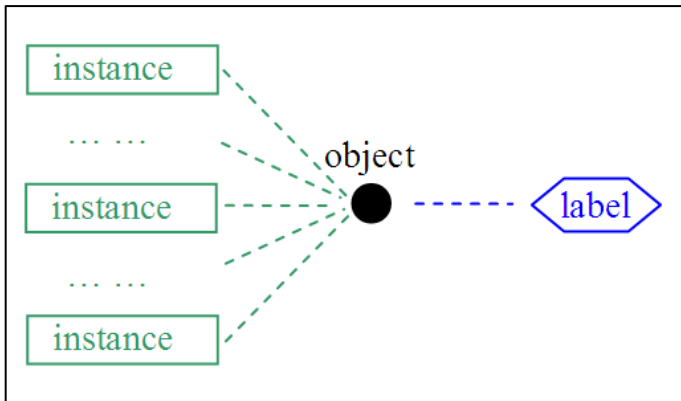
# Why MIIML ? (con't)



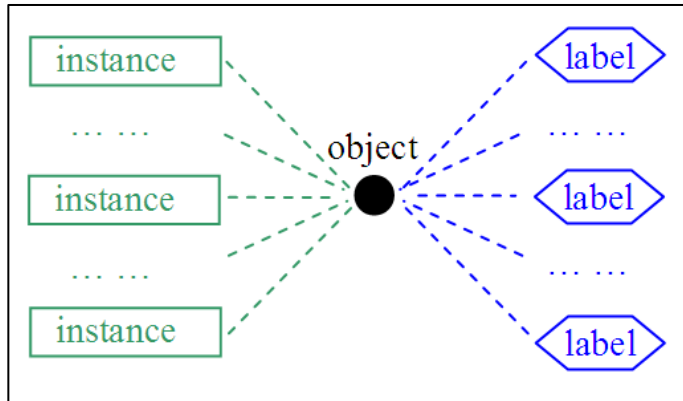
Traditional supervised learning



Multi-label learning



Multi-instance learning



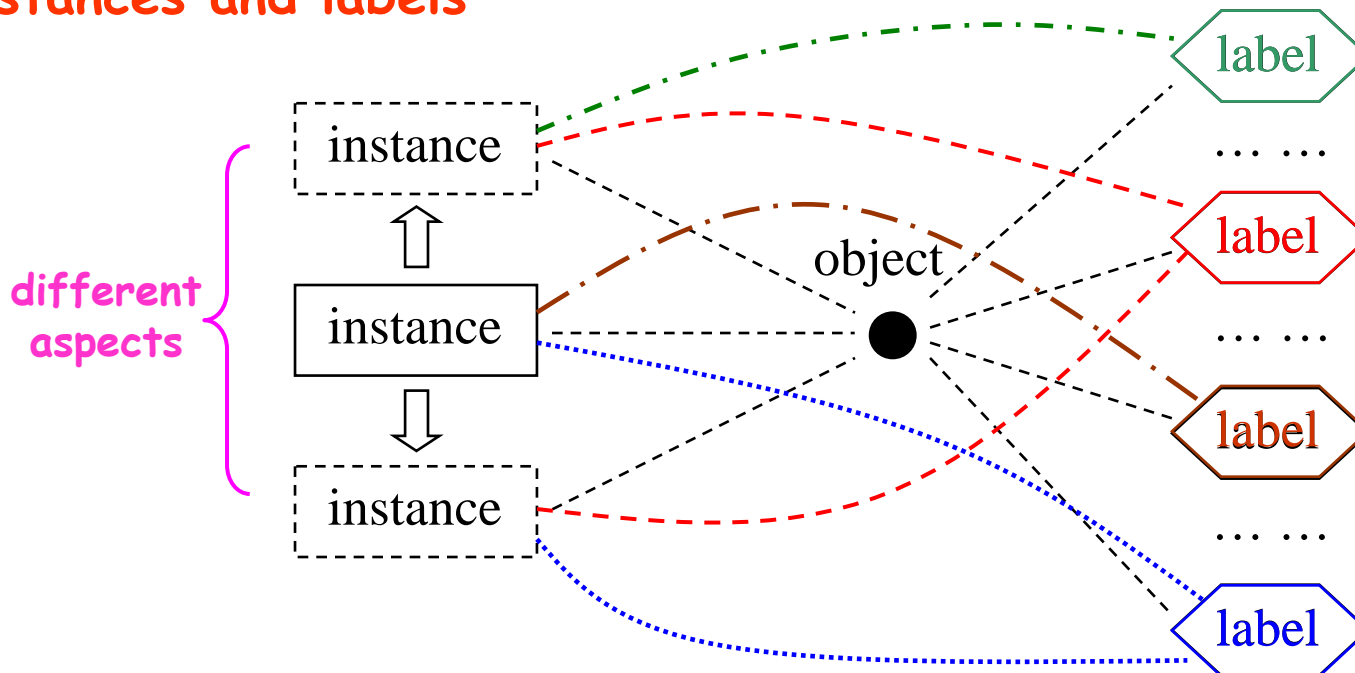
Multi-instance multi-label learning

## Why MIML ? (con't)

To learn an *one-to-many* mapping is an ill-posed problem

Why there are multiple labels?

*many-to-many* mapping seems better; and moreover, **MIML** also offers a possibility for understanding the relationship between instances and labels





## Why MIML ? (con't)

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MIML can also be helpful for learning single-label examples involving complicated high-level concepts

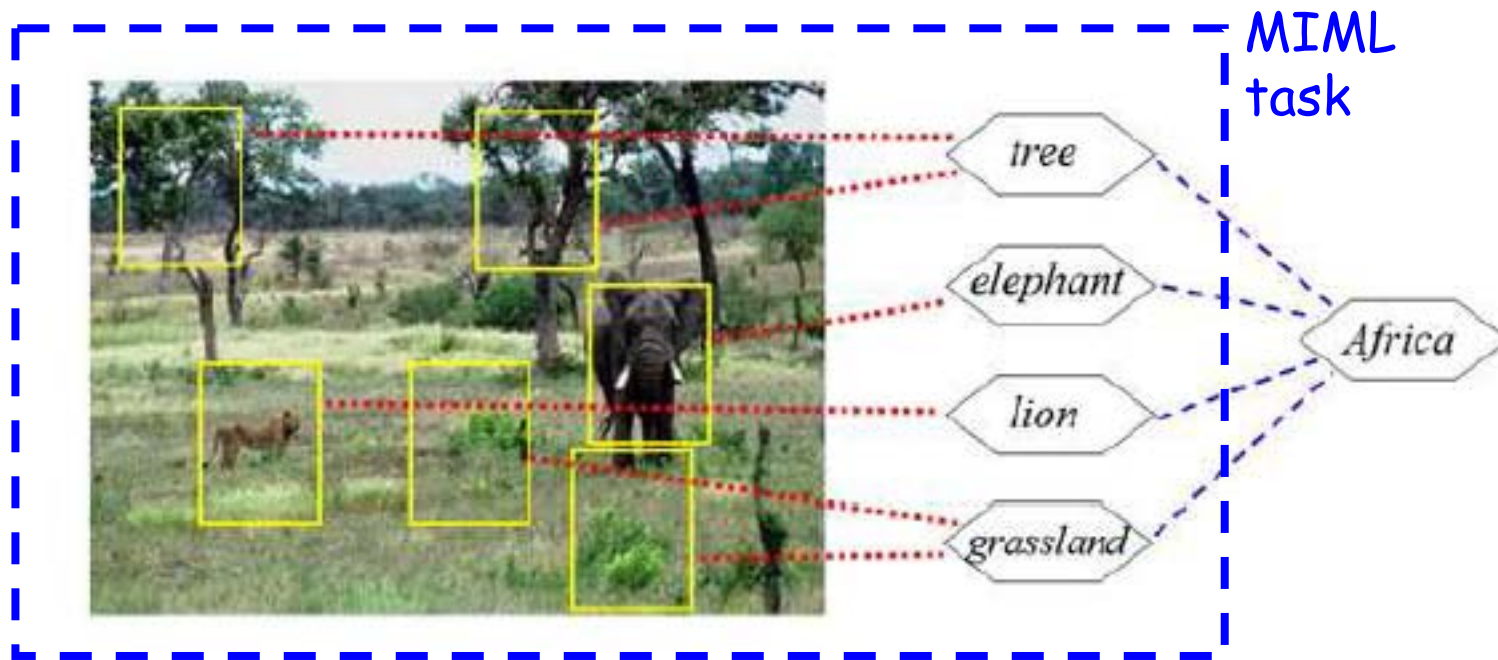


(a) *Africa* is a complicated high-level concept



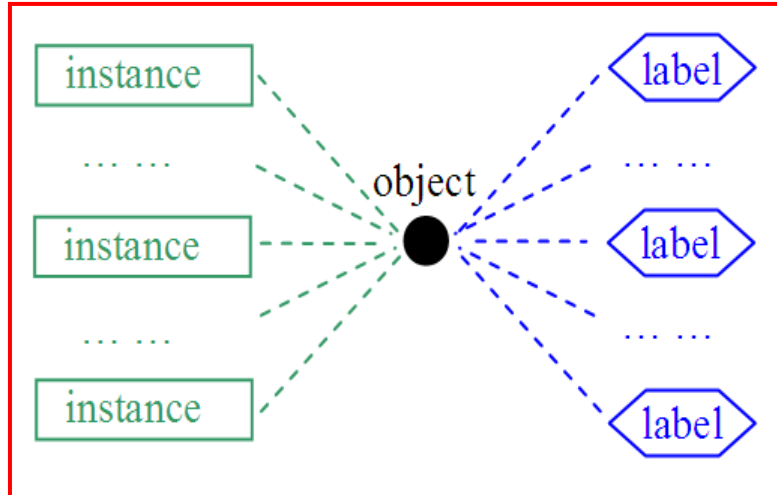
## Why MIML ? (con't)

MIML can also be helpful for learning single-label examples involving complicated high-level concepts



(b) The concept *Africa* may become easier to learn through exploiting some sub-concepts

# Multi-Instance Multi-Label Learning



## MIML task:

To learn a function  $f_{MIML} : 2^{\mathcal{X}} \rightarrow 2^{\mathcal{Y}}$  from a given data set  $\{(X_1, Y_1), (X_2, Y_2), \dots, (X_m, Y_m)\}$ , where  $X_i \subseteq \mathcal{X}$  is a set of instances  $\{\mathbf{x}_1^{(i)}, \mathbf{x}_2^{(i)}, \dots, \mathbf{x}_{n_i}^{(i)}\}$ ,  $\mathbf{x}_j^{(i)} \in \mathcal{X}$  ( $j = 1, 2, \dots, n_i$ ), and  $Y_i \subseteq \mathcal{Y}$  is a set of labels  $\{y_1^{(i)}, y_2^{(i)}, \dots, y_{l_i}^{(i)}\}$ ,  $y_k^{(i)} \in \mathcal{Y}$  ( $k = 1, 2, \dots, l_i$ ).

## MIML:

多示例多标记学习

$\mathcal{X}$  - the instance space

$\mathcal{Y}$  - the set of class labels

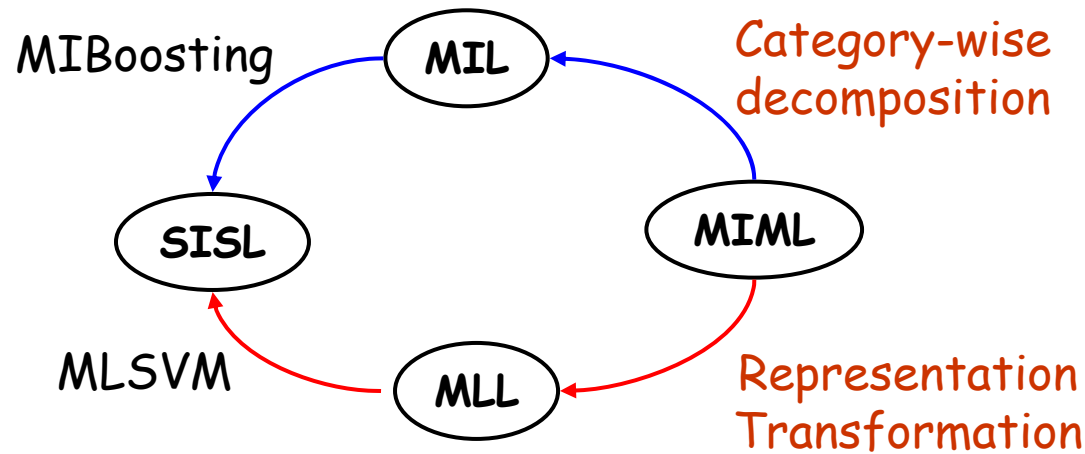
$n_i$  - the number of instances in  $X_i$

$l_i$  - the number of labels in  $Y_i$

# MIMLBoost & MIMLSVM

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## MIMLBoost (an illustration of Solution 1)



## MIMLSVM (an illustration of Solution 2)



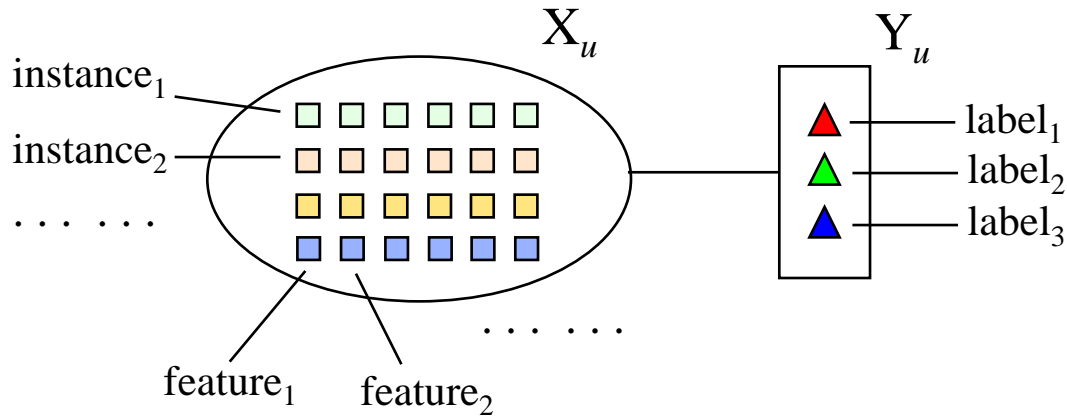
Table 1: The MIMLBOOST algorithm

- 
- 1 Transform each MIML example  $(X_u, Y_u)$  ( $u = 1, 2, \dots, m$ ) into  $|\mathcal{Y}|$  number of multi-instance bags  $\{[(X_u, y_1), \Psi(X_u, y_1)], \dots, [(X_u, y_{|\mathcal{Y}|}), \Psi(X_u, y_{|\mathcal{Y}|})]\}$ . Thus, the original data set is transformed into a multi-instance data set containing  $m \times |\mathcal{Y}|$  number of multi-instance bags, denoted by  $\{[(X^{(i)}, y^{(i)}), \Psi(X^{(i)}, y^{(i)})]\}$  ( $i = 1, 2, \dots, m \times |\mathcal{Y}|$ ).
  - 2 Initialize weight of each bag to  $W^{(i)} = \frac{1}{m \times |\mathcal{Y}|}$  ( $i = 1, 2, \dots, m \times |\mathcal{Y}|$ ).
  - 3 Repeat for  $t = 1, 2, \dots, T$  iterations:
    - 3a Set  $W_j^{(i)} = W^{(i)} / n_i$  ( $i = 1, 2, \dots, m \times |\mathcal{Y}|$ ), assign the bag's label  $\Psi(X^{(i)}, y^{(i)})$  to each of its instances  $(x_j^{(i)}, y^{(i)})$  ( $j = 1, 2, \dots, n_i$ ), and build an instance-level predictor  $h_t[(x_j^{(i)}, y^{(i)})] \in \{-1, +1\}$ .
    - 3b For the  $i$ th bag, compute the error rate  $e^{(i)} \in [0, 1]$  by counting the number of misclassified instances within the bag, i.e.  $e^{(i)} = \frac{\sum_{j=1}^{n_i} [h_t[(x_j^{(i)}, y^{(i)})] \neq \Psi(X^{(i)}, y^{(i)})]}{n_i}$ .
    - 3c If  $e^{(i)} < 0.5$  for all  $i \in \{1, 2, \dots, m \times |\mathcal{Y}|\}$ , go to Step 4.
    - 3d Compute  $c_t = \arg \min_{c_t} \sum_{i=1}^{m \times |\mathcal{Y}|} W^{(i)} \exp[(2e^{(i)} - 1)c_t]$ .
    - 3e If  $c_t \leq 0$ , go to Step 4.
    - 3f Set  $W^{(i)} = W^{(i)} \exp[(2e^{(i)} - 1)c_t]$  ( $i = 1, 2, \dots, m \times |\mathcal{Y}|$ ) and re-normalize such that  $0 \leq W^{(i)} \leq 1$  and  $\sum_{i=1}^{m \times |\mathcal{Y}|} W^{(i)} = 1$ .
  - 4 Return  $Y^* = \{y | \arg_{y \in \mathcal{Y}} \text{sign} \left( \sum_j \sum_t c_t h_t[(x_j^*, y)] \right) = +1\}$  ( $x_j^*$  is  $X^*$ 's  $j$ th instance).
-

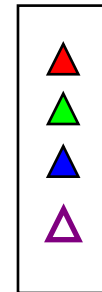
# MIMLBoost

Illustration of the **category-wise decomposition**:

An MIML example  $(X_u, Y_u)$



Label set  $\mathcal{Y}$



# MIMLBoost (con't)

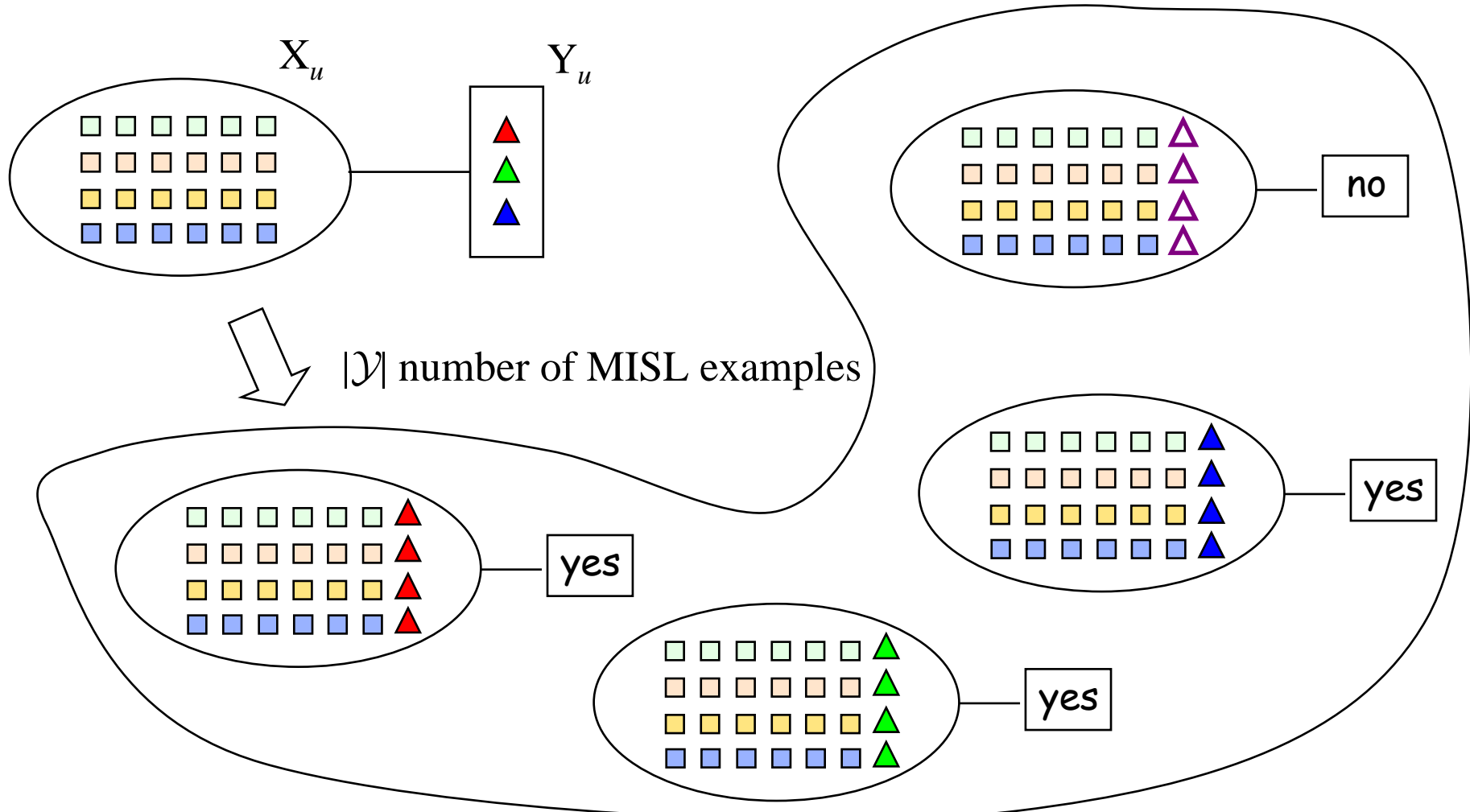


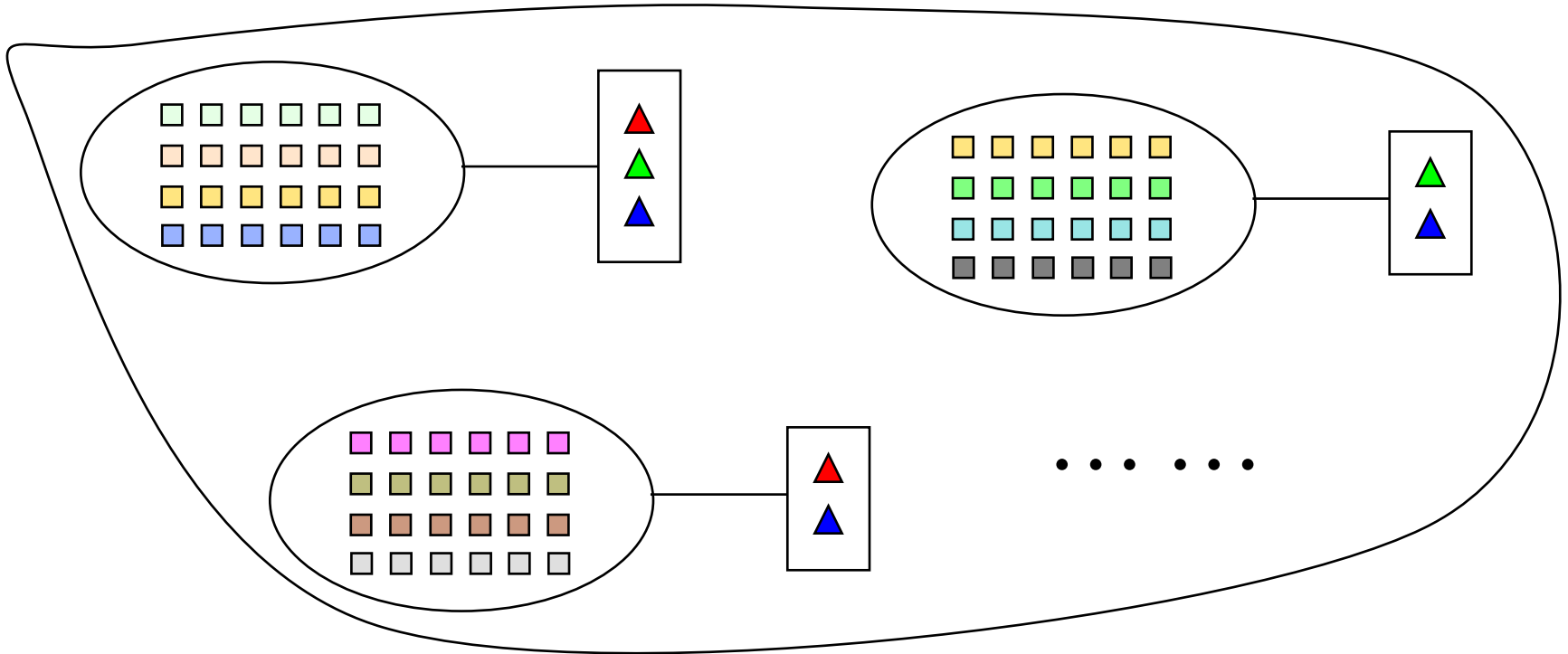
Table 2: The MIMLSVM algorithm

- 
- 1 For MIML examples  $(X_u, Y_u)$  ( $u = 1, 2, \dots, m$ ),  $\Gamma = \{X_u | u = 1, 2, \dots, m\}$ .
  - 2 Randomly select  $k$  elements from  $\Gamma$  to initialize the medoids  $M_t$  ( $t = 1, 2, \dots, k$ ), repeat until all  $M_t$  do not change:
    - 2a  $\Gamma_t = \{M_t\}$  ( $t = 1, 2, \dots, k$ ).
    - 2b Repeat for each  $X_u \in (\Gamma - \{M_t | t = 1, 2, \dots, k\})$ :
 
$$index = \arg \min_{t \in \{1, \dots, k\}} d_H(X_u, M_t), \Gamma_{index} = \Gamma_{index} \cup \{X_u\}.$$
    - 2c  $M_t = \arg \min_{A \in \Gamma_t} \sum_{B \in \Gamma_t} d_H(A, B)$  ( $t = 1, 2, \dots, k$ ).
  - 3 Transform  $(X_u, Y_u)$  into a multi-label example  $(z_u, Y_u)$  ( $u = 1, 2, \dots, m$ ), where  $z_u = (z_{u1}, z_{u2}, \dots, z_{uk}) = (d_H(X_u, M_1), d_H(X_u, M_2), \dots, d_H(X_u, M_k))$ .
  - 4 For each  $y \in \mathcal{Y}$ , derive a data set  $\mathcal{D}_y = \{(z_u, \Phi(z_u, y)) | u = 1, 2, \dots, m\}$ , and then train an SVM  $h_y = SVMTrain(\mathcal{D}_y)$ .
  - 5 Return  $Y^* = \{\arg \max_{y \in \mathcal{Y}} h_y(z^*)\} \cup \{y | h_y(z^*) \geq 0, y \in \mathcal{Y}\}$ , where  $z^* = (d_H(X^*, M_1), d_H(X^*, M_2), \dots, d_H(X^*, M_k))$ .
-

# MIMLSVM

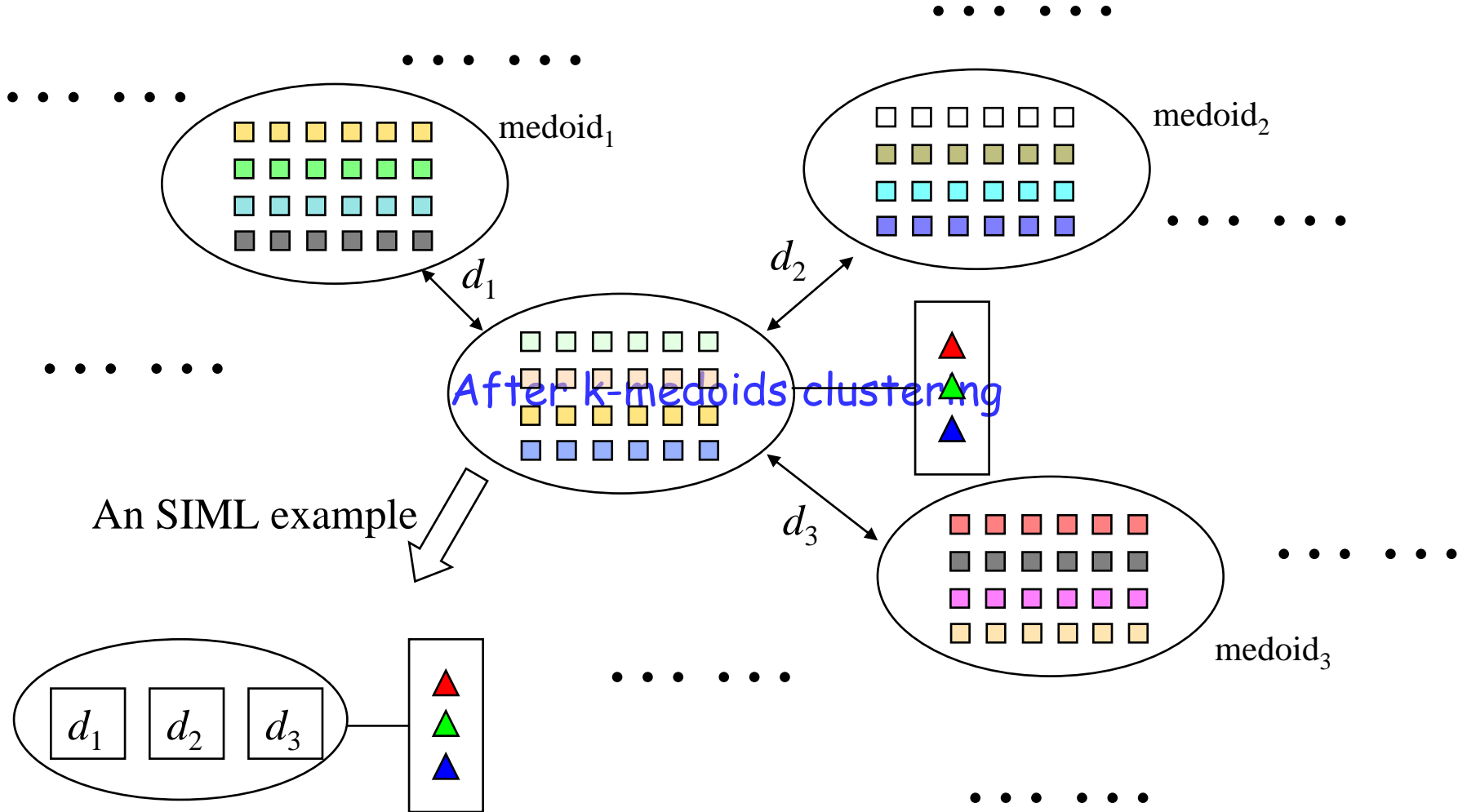
Illustration of the **representation transformation**:

A set of MIML examples



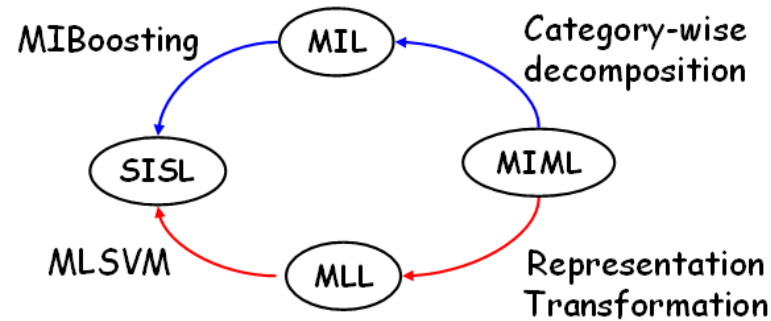


# MIMLSVM (con't)



# Again, Why MIIML?

MIIMLBoost (an illustration of Solution 1)



MIIMLSVM (an illustration of Solution 2)

unambiguous  $\xrightarrow{\hspace{10em}}$  ambiguous

- The MIIML framework incorporates more information (+)
- These solutions degenerate MIIML to solve, while the degeneration loses information (-)

**If (+) > (-), then it is worth doing**

## Scene Classification: Result

Table 3

Results (mean±std.) on scene classification (‘↓’ indicates ‘the smaller the better’; ‘↑’ indicates ‘the larger the better’)

Compared Algorithms	Evaluation Criteria				
	<i>hloss</i> ↓	<i>one-error</i> ↓	<i>coverage</i> ↓	<i>rloss</i> ↓	<i>aveprec</i> ↑
MIMLBOOST	.192±.004	.349±.016	.986±.041	.179±.008	.778±.009
MIMLSVM	.190±.009	.350±.020	1.083±.050	.201±.001	.766±.013
ADTBOOST.MH	.210±.006	.436±.019	1.223±.049	N/A	.718±.012
RANKSVM	.219±.020	.400±.062	1.177±.160	.225±.041	.739±.040
ML- <i>k</i> NN	.191±.006	.370±.017	1.085±.047	.203±.010	.759±.010

The MIML algorithms are apparently superior to non-MIML algorithms

# Text Categorization: Result

Table 4

Results (mean±std.) on text categorization (‘↓’ indicates ‘the smaller the better’; ‘↑’ indicates ‘the larger the better’)

Compared Algorithms	Evaluation Criteria				
	<i>hloss</i> ↓	<i>one-error</i> ↓	<i>coverage</i> ↓	<i>rloss</i> ↓	<i>aveprec</i> ↑
MIMLBOOST	.054±.004	.092±.013	.401±.035	.037±.004	.937±.007
MIMLSVM	.034±.003	.071±.009	.315±.029	.024±.003	.955±.006
ADTBOOST.MH	.055±.004	.120±.016	.409±.046	N/A	.925±.010
RANKSVM	.093±.007	.205±.055	.639±.161	.078±.027	.867±.037
ML- <i>k</i> NN	.067±.005	.191±.017	.683±.052	.085±.008	.871±.010

The MIML algorithms are apparently superior to non-MIML algorithms

# MIML Results

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## Solving MIML problems by degeneration:

- MIMLBoost [Z.-H. Zhou & M.-L. Zhang, NIPS'06]
- MIMLSVM [Z.-H. Zhou & M.-L. Zhang, NIPS'06]

## Solving MIML problems by regularization:

- D-MIMLSVM [Z.-H. Zhou et al., CORR abs/0808.3231]

## Large margin MIML algorithm:

- M3MIML [M.-L. Zhang & Z.-H. Zhou, ICDM'08]

## MIML Results (con't)

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The usefulness of MIML when there is no access to raw objects:

- INSDIF [M.-L. Zhang & Z.-H. Zhou, AAI'07]

MIML to help the learning of complicated high-level concepts:

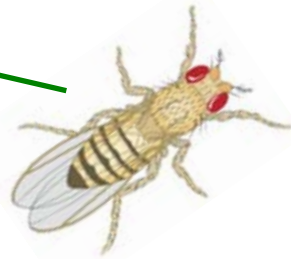
- SUBCOD [Z.-H. Zhou et al., CORR abs/0808.3231]

MIML for image annotation [Z.-J. Zha et al., CVPR'08]

MIML metric learning [R. Jin et al., CVPR'09]

# Drosophila Gene Expression Pattern

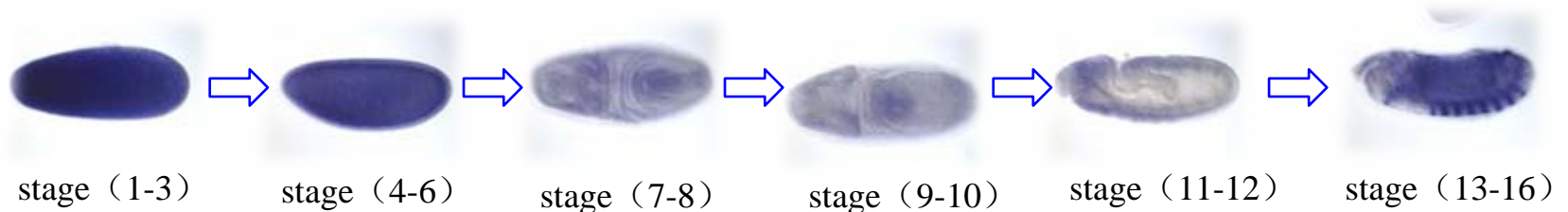
*Drosophila*, or fruit fly, is a model organism widely studied in developmental biology



Gene **RhoGAP71E** expressed stage: 7-8



Gene expression pattern by RNA *in situ* hybridization during *Drosophila* embryogenesis



# The BDGP Project

The *Berkeley Drosophila Genome Project* (BDGP) produced a large amount of spatial-temporal gene expression images



anatomical and developmental ontology terms manually labeled by human curators

Gene: *Actn*



# Difficulty for Automatic Annotation

stage11-12

brain primordium  
visceral muscle primordium  
ventral nerve cord primordium

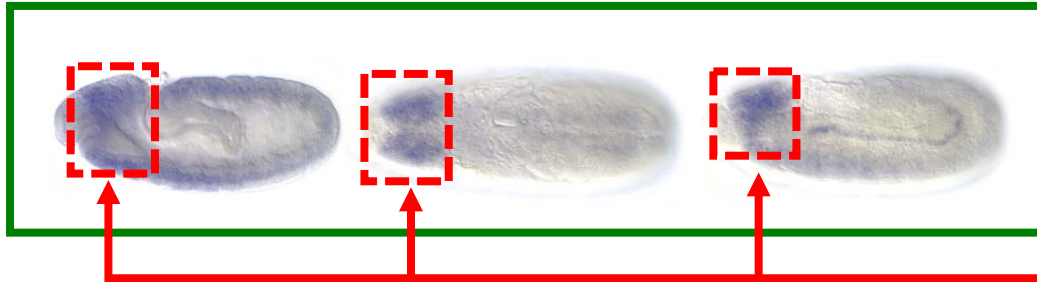
stage13-16

visceral muscle  
embryonic/larval muscle system  
ventral nerve cord  
embryonic brain

brain primordium  
visceral muscle primordium  
ventral nerve cord primordium

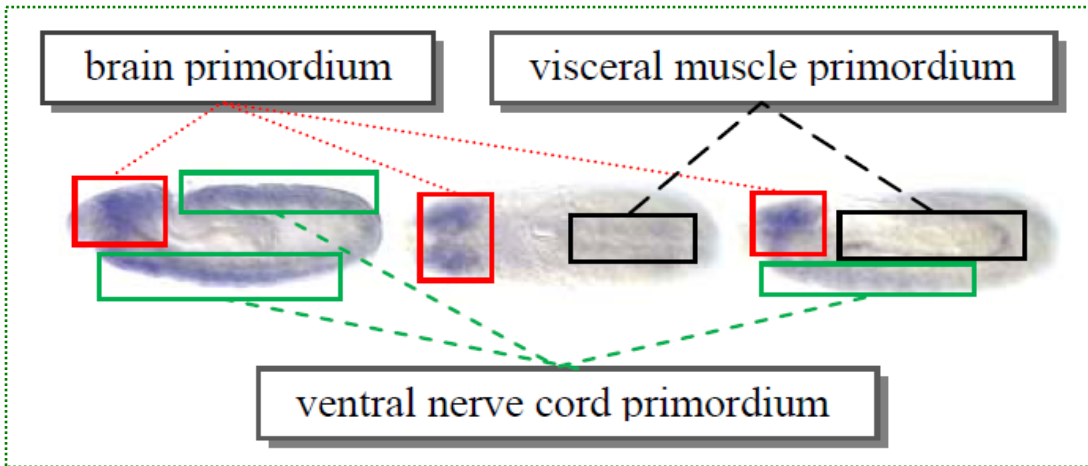
brain primordium  
visceral muscle primordium  
ventral nerve cord primordium

# Difficulty for Automatic Annotation



brain primordium  
visceral muscle primordium  
ventral nerve cord primordium

The terms are body-part related



We do not know  
which term is  
associated with  
which region in  
the images !!

## Generality of the Problem

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A good solution to the *Drosophila* gene expression pattern annotation task will also benefit other bio-problems

e.g., Protein functional prediction

- ✓ many conformations, varying functions
- ✓ Lack knowledge of which conformation is responsible for a specific function

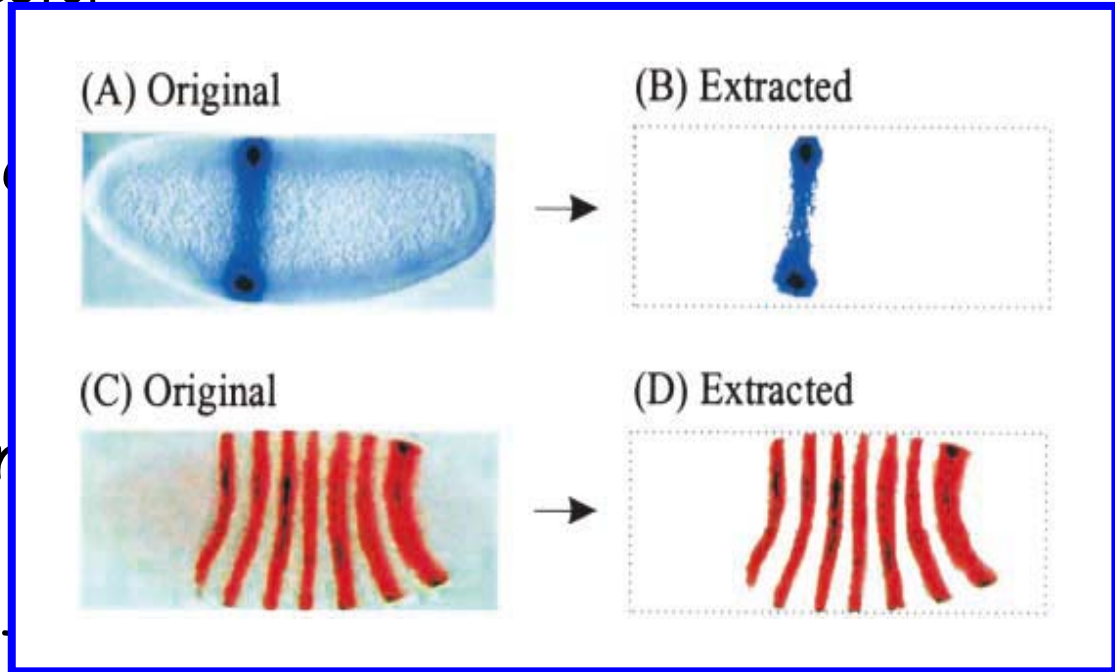
## Previous Solutions

- ✓ **BESTi Algorithm**
  - use images from literatures
  - use binary feature vector

[Kumar et al., Genetics02]

- ✓ **2D Wavelet feature**
  - use BDGP images

- ✓ **Multi-kernel learning**
  - use BDGP images
  - use multi-pyramid ma



## Previous Solutions

✓ **BESTi Algorithm**

[Kumar et al., Genetics02]

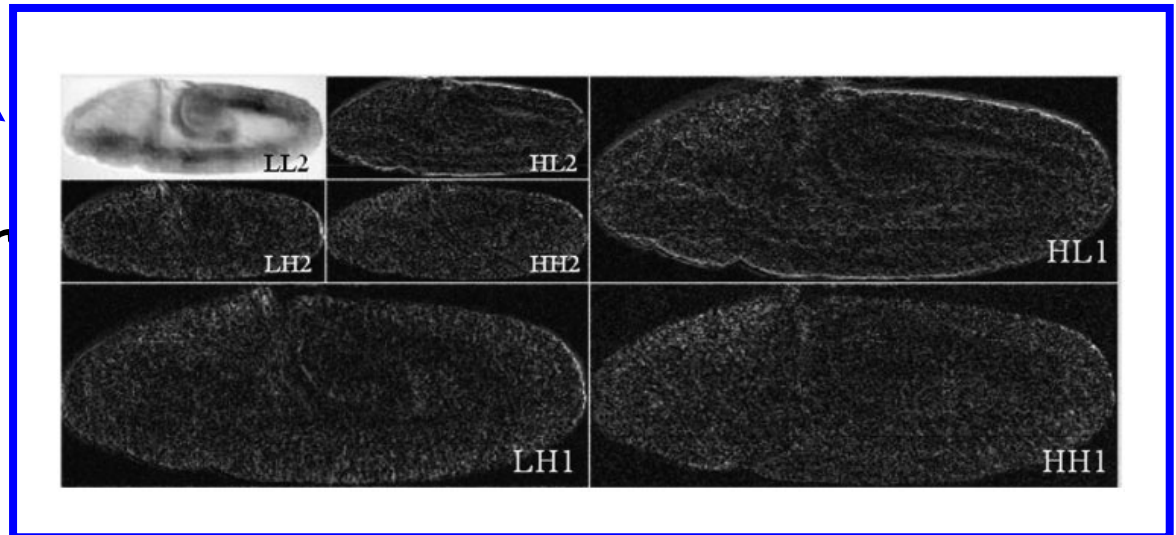
- use images from literatures
- use binary feature vector

✓ **2D Wavelet features, LDA classifier**

- use BDGP images

✓ **Multi-kernel learning**

- use BDGP images
- use multi-pyramid



# Previous Solutions

✓ **BESTI**

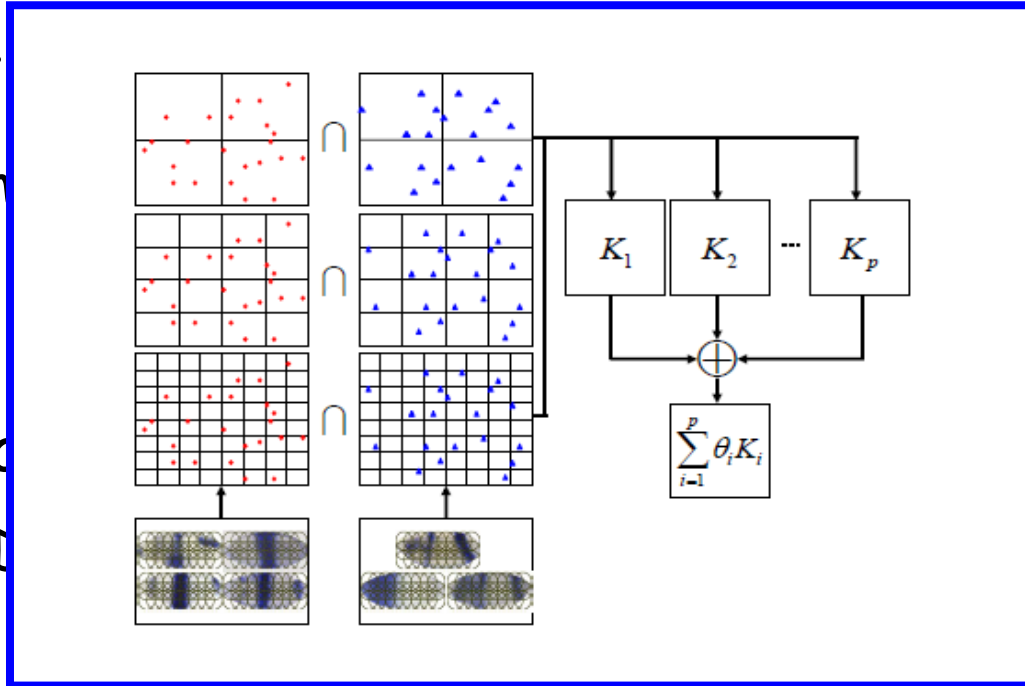
- use im
- use bi

al., Genetics02]

✓ **2D Wo**

- use BD

oinformatics07]

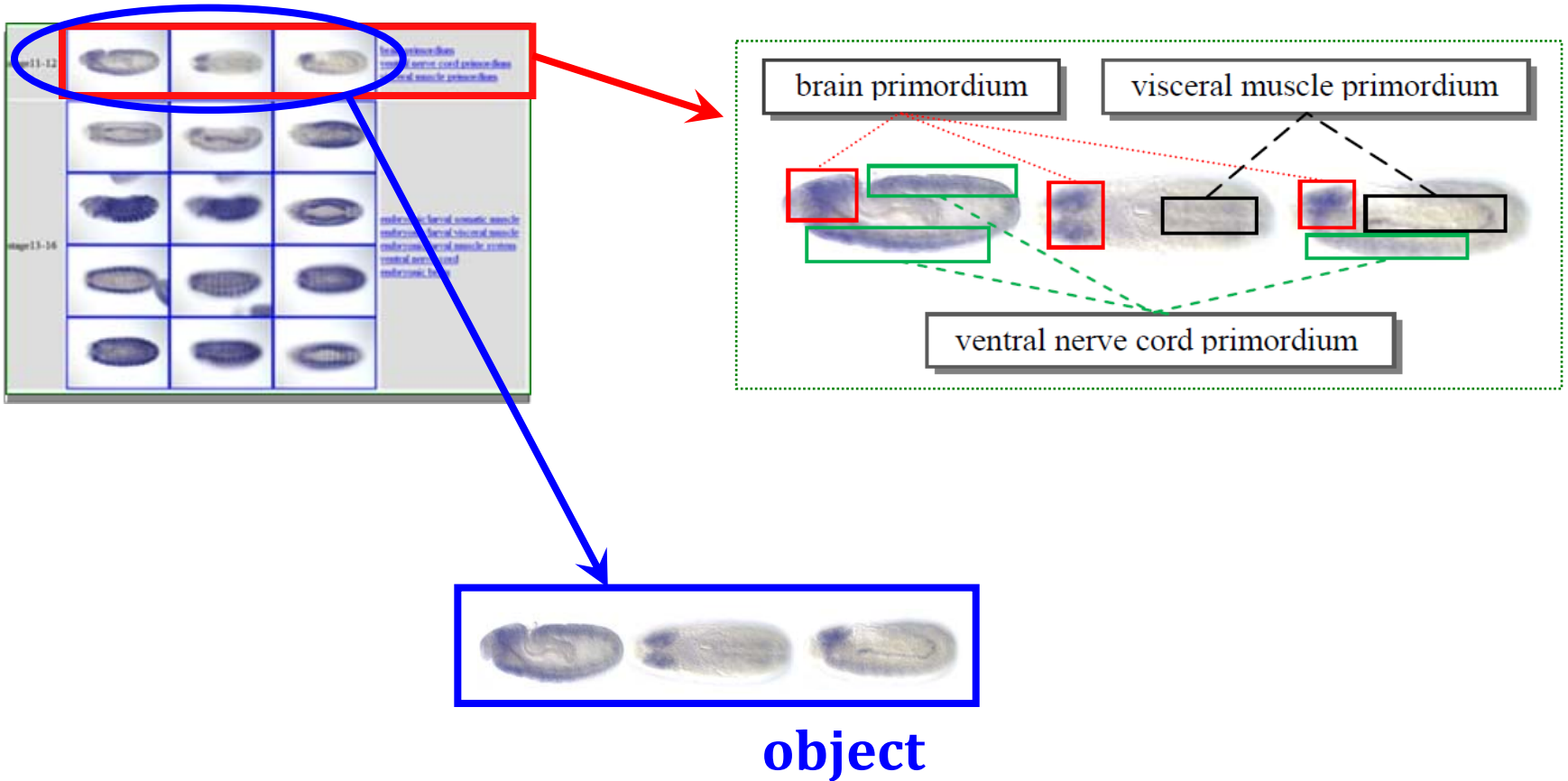


✓ **Multi-kernel learning with hypergraph**

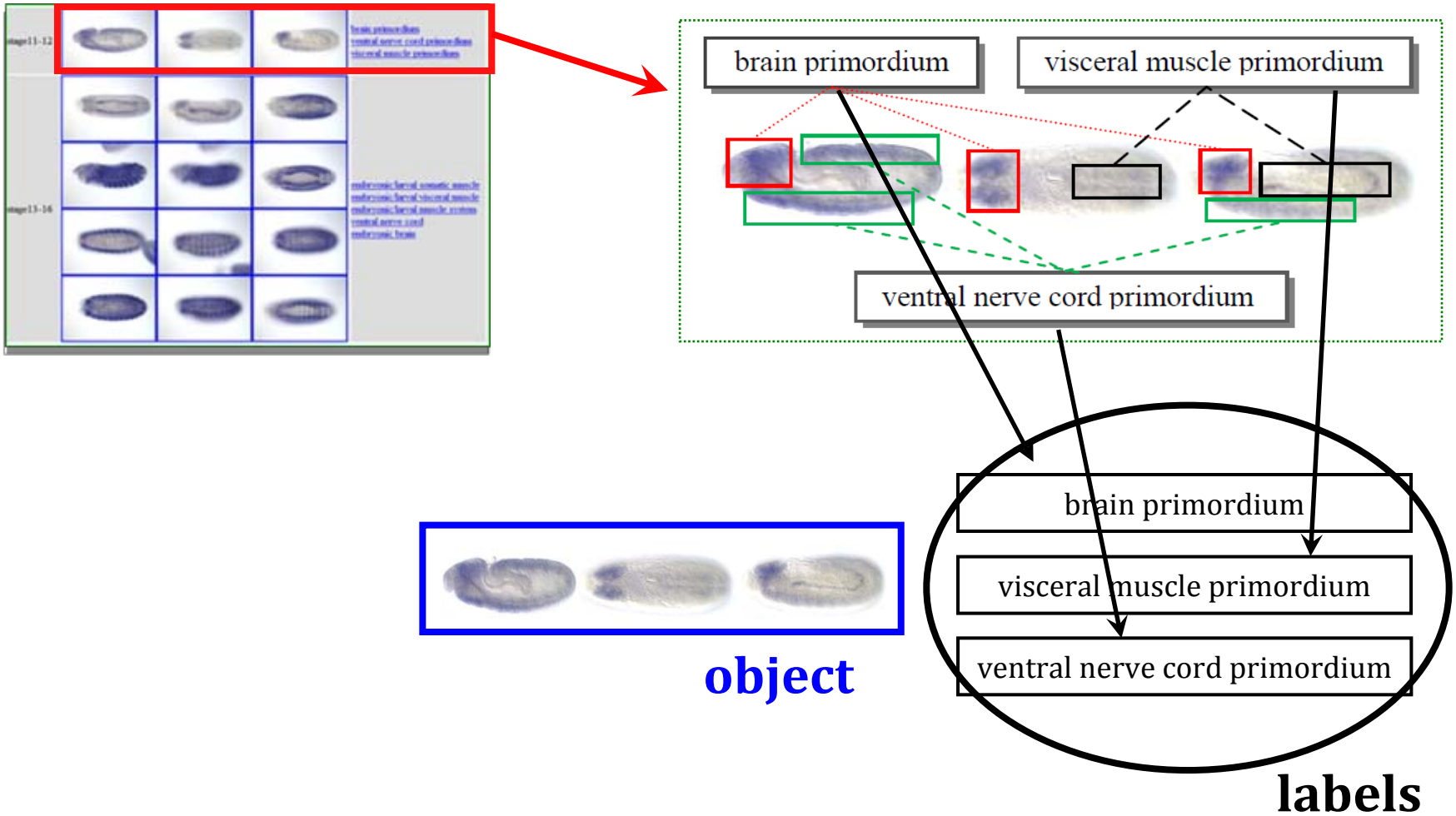
- use BDGP images
- use multi-pyramid match kernel and hypergraph learning

[S. Ji et al., Bioinformatics08]

# Formulated as an MIML Problem

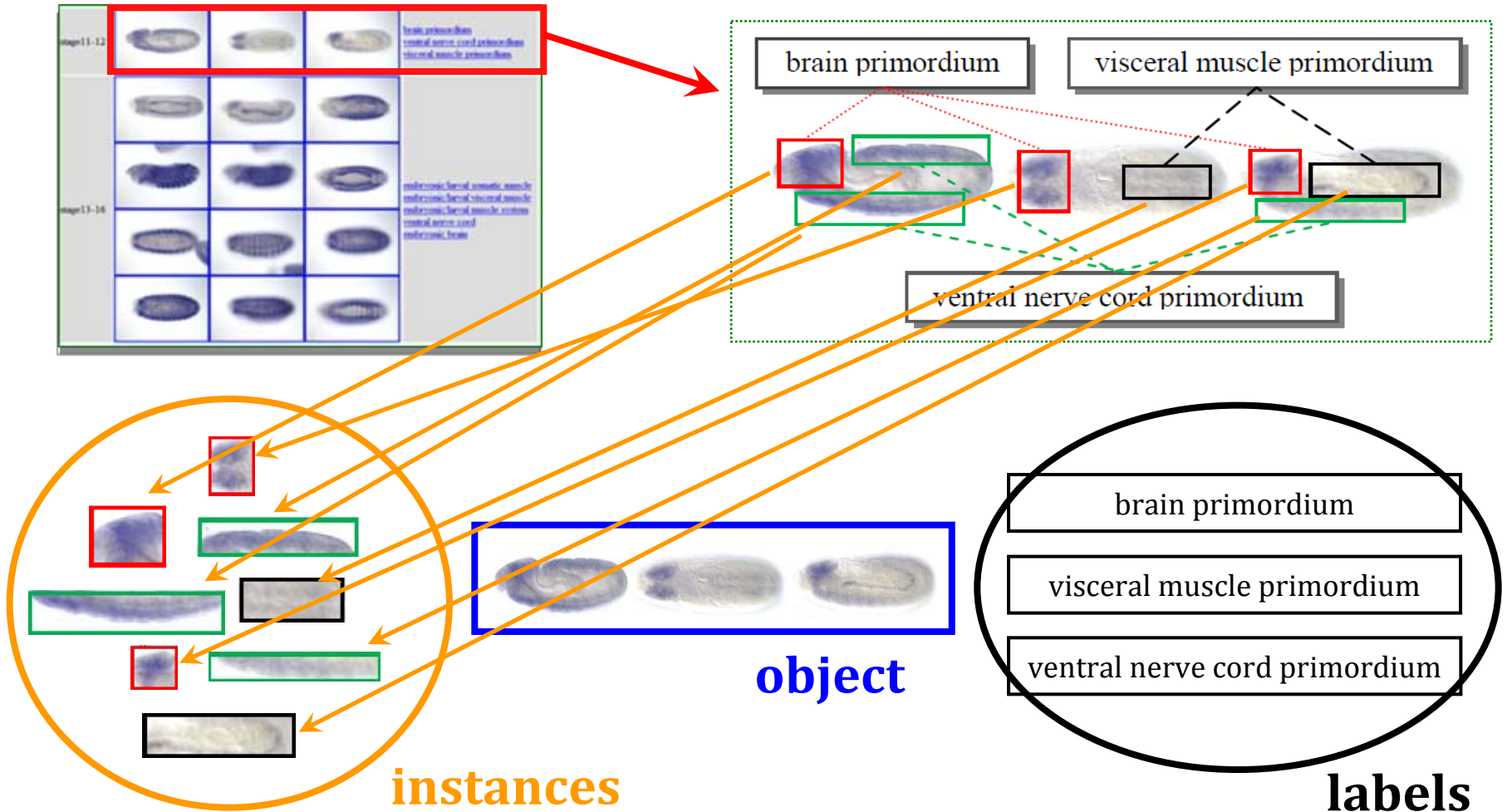


# Formulated as an MIML Problem

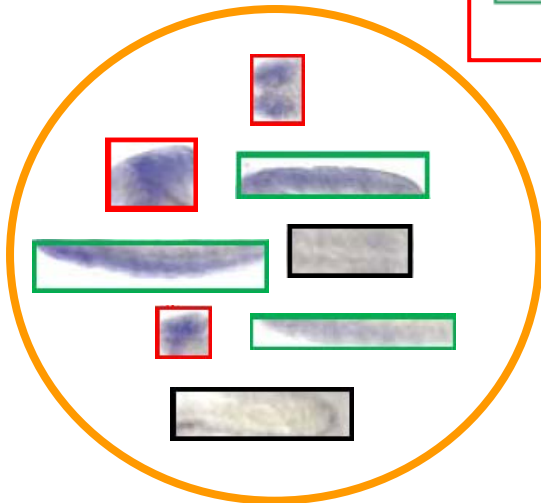
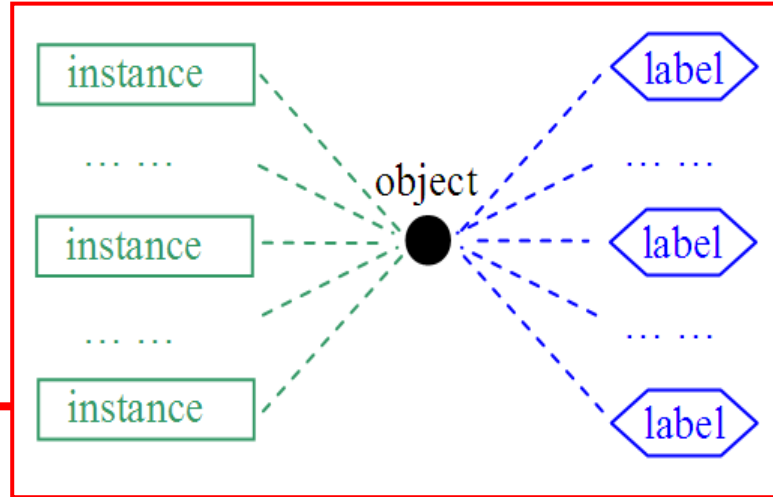




# Formulated as an MIML Problem



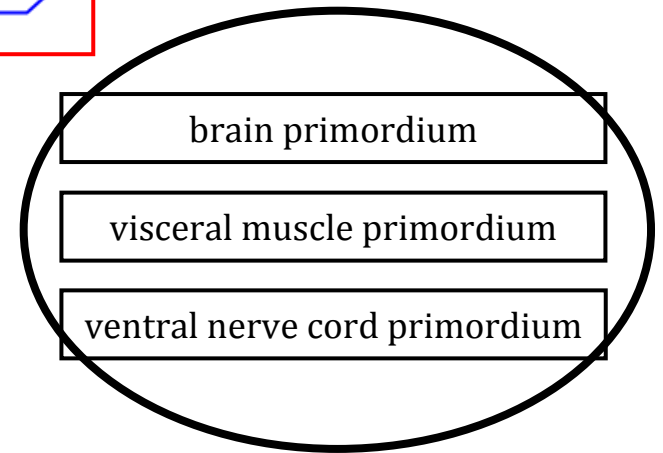
# Formulated as an MIML Problem (con't)



**instances**



**object**



**labels**

# The MIMLSVM+ Algorithm

For each label  $y \in \mathcal{Y}$ , let  $\varphi(X_i, y) = +1$  if  $y \in Y_i$  and -1 otherwise

$$\min_{w, b, \xi} \frac{1}{2} \|w\|^2 + C^+ \sum_{\varphi(X_i, y)=1} \xi_i + C^- \sum_{\varphi(X_i, y)=-1} \xi_i$$

subject to:  $\varphi(X_i, y)(w' \phi(X_i) + b) \geq 1 - \xi_i$   
 $\xi_i \geq 0 \quad (i = 1, 2, \dots, n)$

We set  $C^+ > C^-$  to make the classifier biased toward positive class

# The MIMLSVM+ Algorithm

For each label  $y \in \mathcal{Y}$ , let  $\varphi(X_i, y) = +1$  if  $y \in Y_i$  and -1 otherwise

$$\min_{w, b, \xi} \frac{1}{2} \|w\|^2 + C^+ \sum_{\varphi(X_i, y)=1} \xi_i + C^- \sum_{\varphi(X_i, y)=-1} \xi_i$$

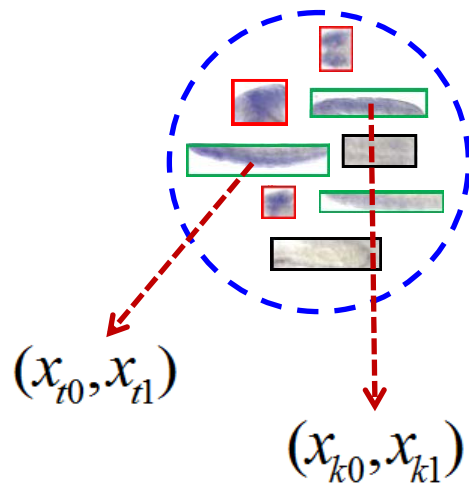
subject to:  $\varphi(X_i, y)(w' \phi(X_i) + b) \geq 1 - \xi_i$   
 $\xi_i \geq 0 \quad (i = 1, 2, \dots, n)$

This involves a kernel function mapping a bag of instances into kernel space. We simply use the *set kernel*:

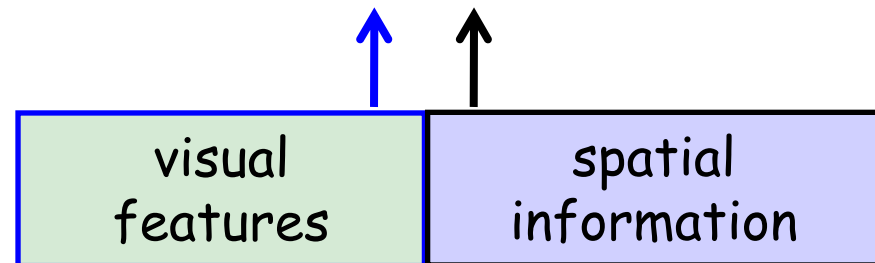
$$K_{SET}(X, X') = \sum_{i=1}^n \sum_{j=1}^m K(x_i, x'_j)$$

## Features Used to Describe Instances

- ✓ visual features of gene expression of patches
- ✓ spatial information of patches



$$X_i = \{x_t\} = \{x_{t0}, x_{t1}\}$$



$$K(x_t, x_k) = e^{-\gamma_1 \|x_{t0} - x_{k0}\|^2 - \gamma_2 \|x_{t1} - x_{k1}\|^2}$$

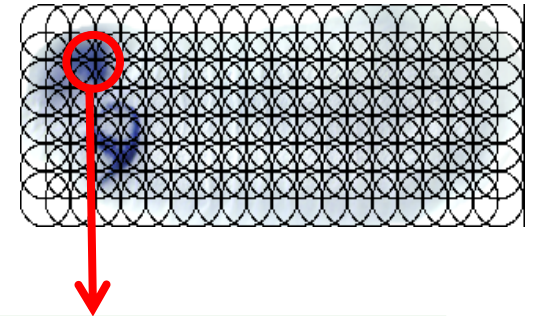
# Experimental Configuration

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## Dataset

**2,816 bags, 2,052,722 instances (15,434 × 133), 119 labels**  
(2,816 image groups, 15,434 images, 133 instances per image, 119 terms)

**Feature** SIFT on dense regular patches  
Center coordinates of patches



**sift & coordinates**

# Evaluation Measures

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## Extended from traditional measures

- ✓ Macro-F1    the larger, the better
- ✓ Micro-F1    the larger, the better
- ✓ AUC (Area under ROC curve)    the larger, the better

## Multi-Label measures

- ✓ Average precision    the larger, the better
- ✓ One-error    the smaller, the better
- ✓ Coverage    the smaller, the better
- ✓ Ranking loss    the smaller, the better
- ✓ Hamming loss    the smaller, the better

# Compared Methods

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## Existing methods

- ✓ MKL-PMK [S. Ji et al., Bioinformatics08]
- ✓ MIML-SVM [Z.-H. Zhou and M.-L. Zhang, NIPS'06]

## Degenerated variants of MIMLSVM<sup>+</sup>

- ✓ MIML-SVM<sub>sv</sub><sup>+</sup> : Concatenate visual and spatial information
- ✓ MIML-SVM<sub>v</sub><sup>+</sup> : Use only visual features



# Experimental Results

50% train 50% test, 30 runs with random partitions

# terms	# groups	Algorithms	macro-F1 ↑	micro-F1 ↑	AUC ↑	Ave. Precision ↑	one-error ↓	coverage ↓	Rankloss ↓	Hammloss ↓
10	222	MIMLSVM <sup>+</sup>	0.643±0.011	0.689±0.007	0.883±0.004	0.779±0.005	0.272±0.008	2.994±0.056	0.150±0.006	0.150±0.004
		MIMLSVM <sup>+</sup> <sub>SV</sub>	0.627±0.010	0.676±0.006	0.869±0.004	0.773±0.005	0.277±0.011	3.073±0.048	0.157±0.004	0.156±0.003
		MIMLSVM <sup>+</sup> <sub>V</sub>	0.619±0.011	0.667±0.007	0.863±0.004	0.764±0.005	0.291±0.009	3.139±0.044	0.164±0.004	0.160±0.003
		MKL-PMK	0.584±0.009	0.621±0.009	0.825±0.006	0.722±0.007	0.343±0.011	3.483±0.072	0.198±0.006	0.196±0.006
20	247	MIMLSVM <sup>+</sup>	0.468±0.015	0.587±0.007	0.862±0.003	0.673±0.008	0.357±0.011	6.189±0.117	0.152±0.005	0.114±0.002
		MIMLSVM <sup>+</sup> <sub>SV</sub>	0.454±0.012	0.574±0.008	0.845±0.003	0.660±0.009	0.364±0.013	6.481±0.119	0.163±0.005	0.118±0.003
		MIMLSVM <sup>+</sup> <sub>V</sub>	0.445±0.012	0.566±0.006	0.840±0.004	0.651±0.008	0.377±0.011	6.609±0.114	0.169±0.004	0.119±0.002
		MKL-PMK	0.410±0.007	0.506±0.006	0.771±0.006	0.580±0.007	0.445±0.009	8.082±0.122	0.230±0.005	0.144±0.003
30	2646	MIMLSVM <sup>+</sup>	0.368±0.012	0.541±0.007	0.850±0.003	0.623±0.007	0.377±0.010	9.406±0.173	0.153±0.003	0.087±0.002
		MIMLSVM <sup>+</sup> <sub>SV</sub>	0.354±0.001	0.527±0.006	0.829±0.004	0.605±0.007	0.388±0.010	9.964±0.195	0.166±0.004	0.090±0.002
		MIMLSVM <sup>+</sup> <sub>V</sub>	0.340±0.012	0.517±0.007	0.822±0.004	0.596±0.007	0.399±0.010	10.183±0.189	0.171±0.004	0.091±0.002
		MKL-PMK	0.310±0.008	0.455±0.008	0.741±0.007	0.511±0.008	0.488±0.011	13.010±0.2413	0.243±0.006	0.142±0.003

**MIMLSVM+ achieves the best performance on ALL cases and ALL evaluation measures**

## Experimental Results (con't)

Since MIMLSVM could not work on the previous large data sets, we extract a smaller data set via random sampling

**167 bags, 57,323 instances (431 × 133), 10 labels**

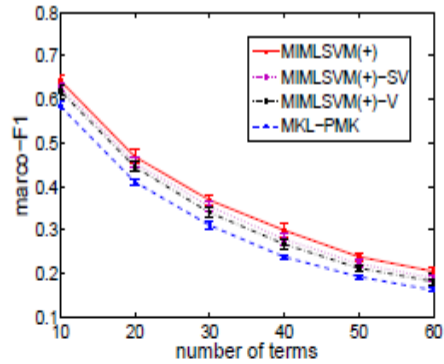
(167 image groups, 431 images, 133 inst per image, 10 terms)

20 runs with random splits of training/test sets

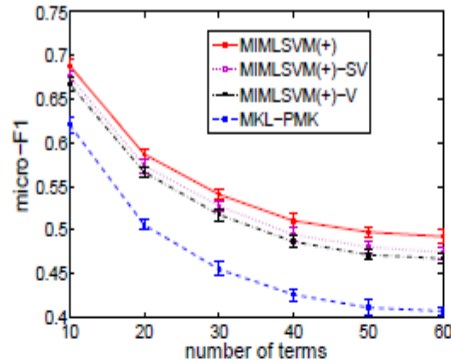
# terms	# groups	Algorithms	macro-F1 ↑	micro-F1 ↑	AUC ↑	Ave. Precision ↑	one-error ↓	coverage ↓	Rankloss ↓	Hammloss ↓
10	167	MIMLSVM <sup>+</sup>	0.460±0.041	0.606±0.026	0.807±0.191	0.733±0.019	0.311±0.034	3.508±0.262	0.186±0.015	0.171±0.019
		MIMLSVM <sup>+</sup> <sub>SV</sub>	0.424±0.049	0.569±0.033	0.774±0.017	0.710±0.027	0.354±0.047	3.667±0.199	0.204±0.016	0.191±0.015
		MIMLSVM	0.176±0.047	0.367±0.054	0.629±0.041	0.592±0.028	0.468±0.060	4.792±0.300	0.318±0.029	0.241±0.097

→ **MIMLSVM+ achieves the best performance on ALL evaluation measures**

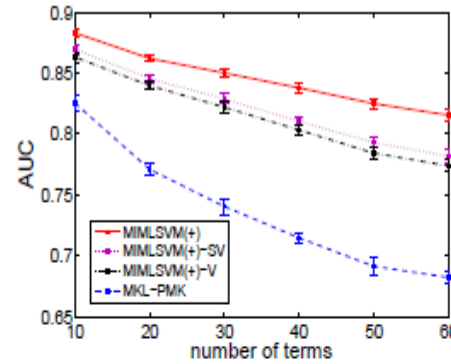
# Experimental Results (con't)



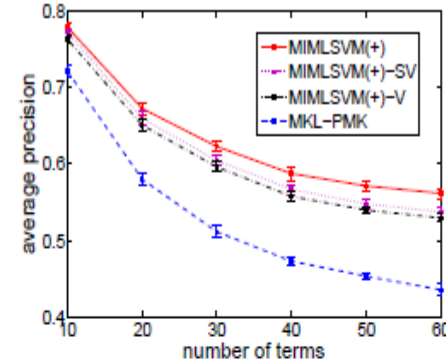
(a) *macro-F1* ↑



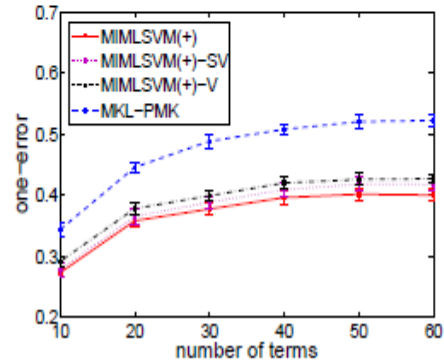
(b) *micro-F1* ↑



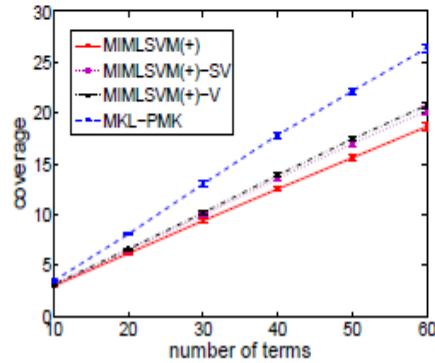
(c) *AUC* ↑



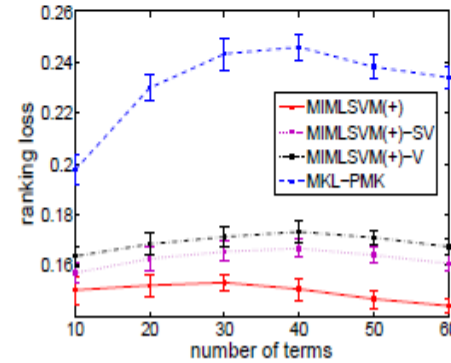
(d) *average precision* ↑



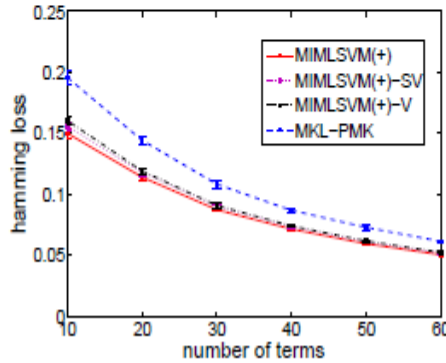
(e) *one-error* ↓



(f) *coverage* ↓



(g) *ranking loss* ↓



(h) *hamming loss* ↓

The comparison under different number of labels (annotation terms)

## MIML Papers

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- ✓ Y.-X. Li, S. Ji, J. Ye, S. Kumar, and Z.-H. Zhou. Drosophila gene expression pattern annotation through multi-instance multi-label learning. [IJCAI'09](#)
- ✓ S. Wang, R. Jin, and Z.-H. Zhou. Learn a distance metric from multi-instance multi-label data. [CVPR'09](#)
- ✓ M.-L. Zhang and Z.-H. Zhou. M<sup>3</sup>MIML: A maximum margin method for multi-instance multi-label learning. [ICDM'08](#)
- ✓ Z.-H. Zhou, M.-L. Zhang, S.-J. Huang, Y.-F. Li. MIML: A Framework for Learning with Ambiguous Objects. [CORR abs/0808.3231](#)
- ✓ M.-L. Zhang, Z.-H. Zhou. Multi-label learning by instance differentiation. [AAAI'07](#), pp.669-674
- ✓ Z.-H. Zhou, M.-L. Zhang. Multi-instance multi-label learning with application to scene classification. [NIPS'06](#), pp.1609-1616.

# MIML Resources

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## Codes:

- MIMLBoost & MIMLSVM:

<http://cs.nju.edu.cn/zhouzh/zhouzh.files/publication/annex/MIMLBoost&MIMLSVM.htm>

- InsDif:

<http://cs.nju.edu.cn/zhouzh/zhouzh.files/publication/annex/InsDif.htm>

- M<sup>3</sup>MIML:

<http://cs.nju.edu.cn/zhouzh/zhouzh.files/publication/annex/M3MIML.htm>

## Data:

- <http://cs.nju.edu.cn/zhouzh/zhouzh.files/publication/annex/miml-image-data.htm>
- <http://cs.nju.edu.cn/zhouzh/zhouzh.files/publication/annex/miml-text-data.htm>

# Thanks!





# 机器学习周(南京大学)

## Machine Learning Week (Nanjing University)



**ACML2009**

**Nov.2-4**

### The 1st Asian Conference on Machine Learning

The 1st Asian Conference on Machine Learning (ACML'09) aims at providing a leading international forum for researchers in machine learning and related fields to share their new ideas and achievements.

*Keynote speaker:*

Thomas Dietterich

(Oregon State University; Founding Chair of IMLS)

*Invited speaker:*

Masashi Sugiyama

(Tokyo Institute of Technology)

Qiang Yang

(Hong Kong University of Science and Technology)

Registration and other details please visit:

<http://lamda.nju.edu.cn/conf/acml09>

(ACML2009将为中国大陆学生提供低价旁听注册, 详情请见会议网站)



**MLA2009**

**11月6-8日**

### 第七届机器学习及其应用研讨会

第七届“机器学习及其应用”研讨会由南京大学计算机软件新技术国家重点实验室主办, 将于2009年11月6-8日在南京举行。该研讨会拟邀请海内外活跃在机器学习及相关领域研究第一线的10余位华人学者前来做特邀报告, 研讨会仍将举行Top Conference Review (“过去一年顶级会议回顾”)以及墙展交流等活动。

按惯例, 该研讨会不征文、不收费, 欢迎机器学习及相关领域的学者、研究生前来旁听特邀报告并参加讨论。

具体情况请见: <http://lamda.nju.edu.cn/conf/mla09>



**SSMLA2009**

**11月6-8日**

### 第四届机器学习及其应用学生研讨会

为了促进机器学习及相关领域的研究生之间以及研究生与资深学者之间的交流, 在第7届机器学习及其应用研讨会期间, 将举行第4届机器学习及其应用学生研讨会(SSMLA'09)。SSMLA'09以Poster(墙展张贴)的方式进行。除部分特邀Poster外, SSMLA'09将公开征集学生Poster, 通过评审的学生Poster将获得到场展示的机会。会议还将评出最佳学生Poster奖。

具体情况请见: <http://lamda.nju.edu.cn/conf/ssmla09>

