

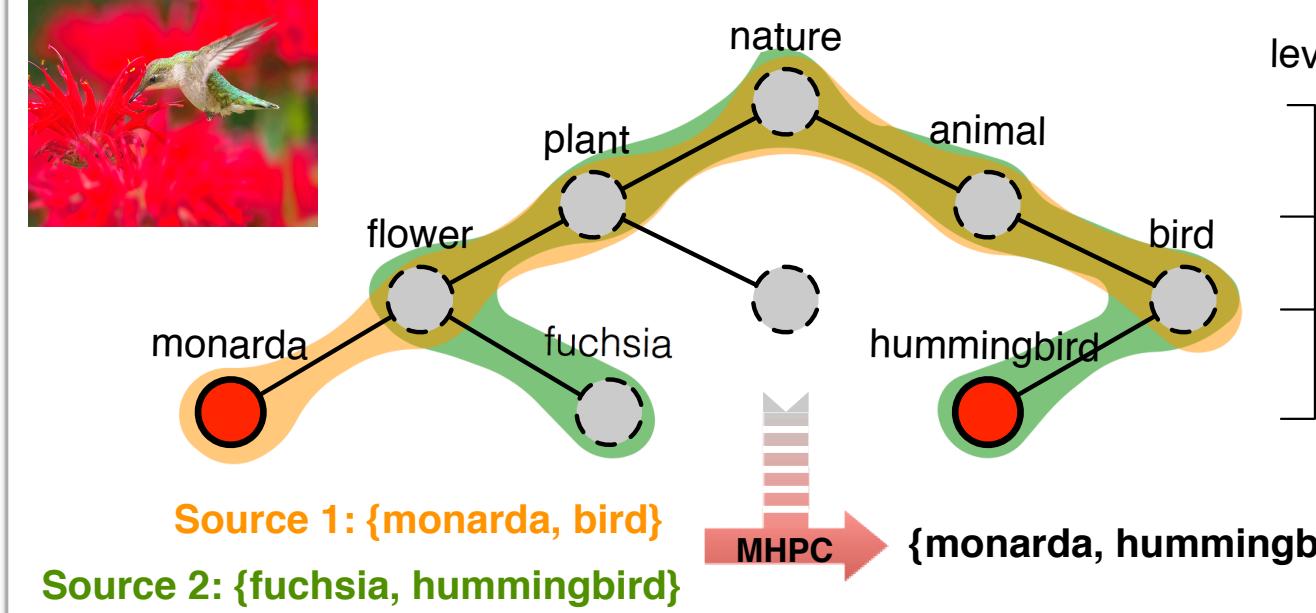
# Multi-source Hierarchical Prediction Consolidation

Chenwei Zhang<sup>1</sup> Sihong Xie<sup>2</sup> Yaliang Li<sup>3</sup> Jing Gao<sup>3</sup> Wei Fan<sup>4</sup> Philip S. Yu<sup>1,5</sup>

<sup>1</sup>University of Illinois at Chicago <sup>2</sup>Lehigh University <sup>3</sup>SUNY Buffalo <sup>4</sup>Baidu Research Big Data Lab <sup>5</sup>Tsinghua University



## 1. Multi-source Hierarchical Prediction Consolidation Problem



**Multiple information sources** may provide labeling information on the same instance simultaneously.

**Consolidation on the output-level:** raw features are neglected or withheld due to storage limitation or privacy concerns.

**Label hierarchies** are prevalently observed.

Traditionally, only “flat” labels are considered.

**Problem Studied:** incorporate the **label hierarchy** into the prediction consolidation process when **only the labeling information** from **multiple information sources** are available.

**Challenges:** label vagueness, label ambiguity, label sparsity.

## 2. Modeling Hierarchical Consensus

### Minimizing the Consensus Cost

An optimization function to minimize the consensus cost among information sources

$$\min_{\bar{Y}} \frac{1}{M} \sum_{m=1}^M \|\hat{Y} - Y_m\|_F^2 + \lambda \sum_{i=1}^N \sum_{j=1}^N W_{ij} \|\hat{Y}_{(i)} - \hat{Y}_{(j)}\|_2^2$$

Favors the smoothness of the consolidation result from all the information sources

Regularizes the model to ensure hierarchical label constraints

A closed-form solution:

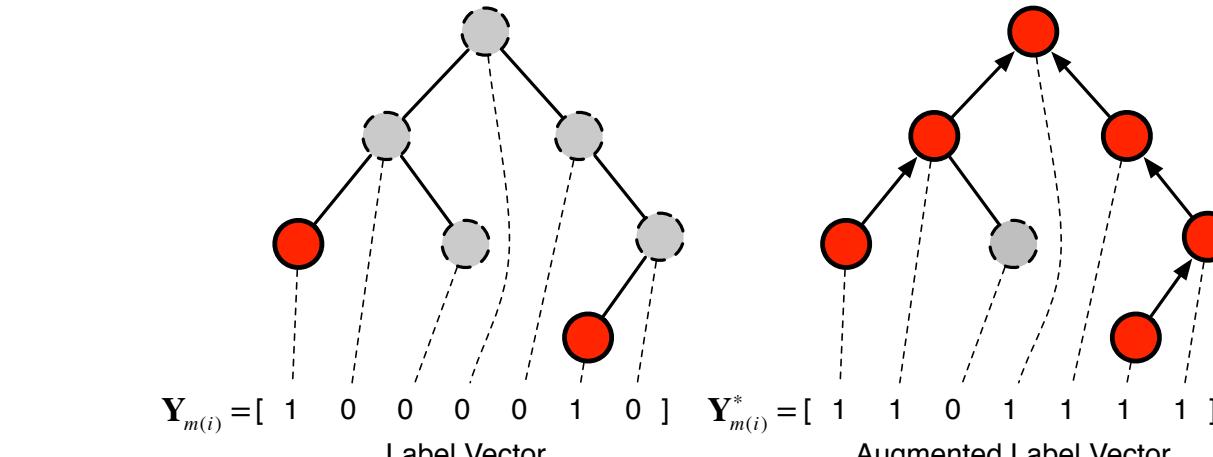
$$\hat{Y} = \left(1 + \lambda \cdot (\mathbf{D}^{-\frac{1}{2}}(\mathbf{D} - \mathbf{W})\mathbf{D}^{-\frac{1}{2}})\right)^{-1} \frac{1}{M} \sum_{m=1}^M \hat{Y}_m = (1 + \lambda \cdot \mathbf{L})^{-1} \bar{Y}$$

$\mathbf{L}$  regularizes the simple averaging result and guides the  $\bar{Y}$  towards a global consensus  $\hat{Y}$ .

### Estimating the Hierarchical Similarity

$$W_{ij} = \exp\left(-\frac{1}{\sigma} \sqrt{\sum_{k=1}^K \vec{S}_k (\hat{Y}_{(i)}^k - \hat{Y}_{(j)}^k)^2}\right)$$

Label augmentation based on the label Hierarchy:



The label occurrence in augmented label vectors:

$$\bar{C}_k = \sum_{m=1}^M \sum_{i=1}^N Y_{m(i)}^{k*} \quad \vec{S}_k = B\left(\frac{\alpha}{2}; \bar{C}_k, N - \bar{C}_k + 1\right)$$

Uncertainty modeled by the Beta distribution:

$$B\left(\frac{\alpha}{2}; \bar{C}_k, N - \bar{C}_k + 1\right) < \theta_k < B\left(1 - \frac{\alpha}{2}; \bar{C}_k, N - \bar{C}_k + 1\right)$$

## 5. Acknowledgements

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## 3. Consolidating Hierarchical Labels

The MHPC Algorithm: a totally **unsupervised**, **iterative** updating algorithm.

### Minimizing the Consensus Cost

$$\begin{aligned} (\hat{Y})_{t_{x+1}} &= \left(1 + \lambda \cdot (\mathbf{L})_{t_x}\right)^{-1} (\hat{Y})_{t_x} \\ \hat{Y} &= (1 + \lambda \cdot \mathbf{L})^{-1} \bar{Y} \\ \text{Initialize: } (\hat{Y})_{t_0} &= \bar{Y} \end{aligned}$$

accumulate the consolidation progresses from previous iterations.

The two-phase updating schema terminates when updates on the consolidation result  $(\hat{Y})_{t_{x+1}}$  is no longer significant after an iteration.

### Estimating the Hierarchical Similarity

$$(W_{ij})_{t_{x+1}} = \exp\left(-\frac{1}{\sigma} \sqrt{\sum_{k=1}^K \vec{S}_k ((\hat{Y}_{(i)})_{t_x} - (\hat{Y}_{(j)})_{t_x})^2}\right)$$

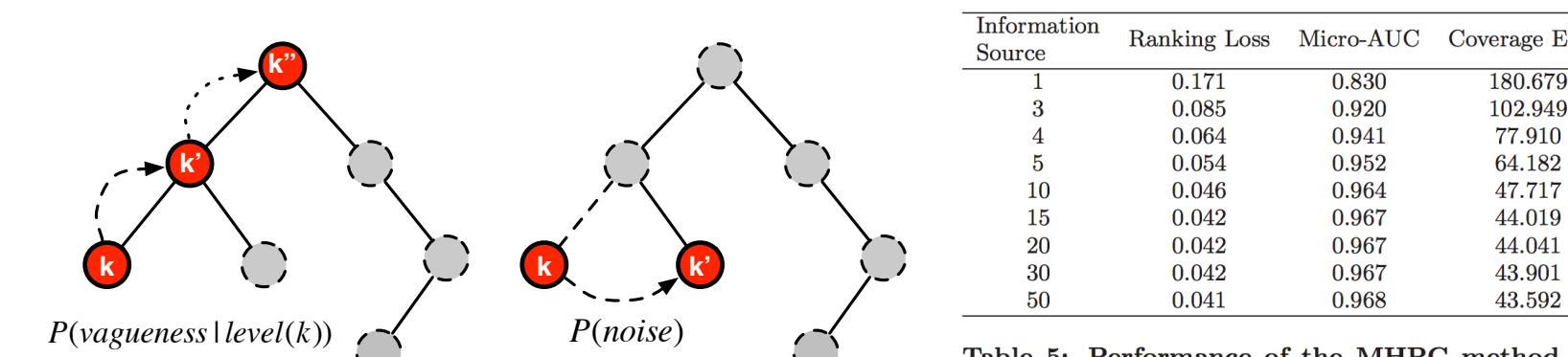
Initialize:

$$(W_{ij})_{t_0} = \exp\left(-\frac{1}{\sigma} \sqrt{\sum_{k=1}^K \vec{S}_k (\bar{Y}_{(i)}^k - \bar{Y}_{(j)}^k)^2}\right)$$

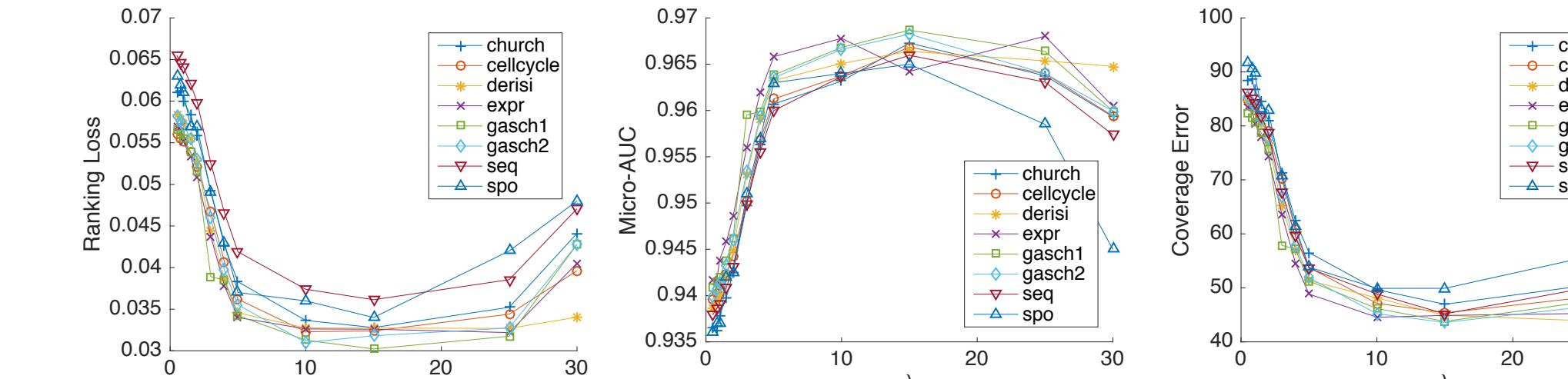
## 4. Experiments

### Multi-source Yeast Genome Annotation

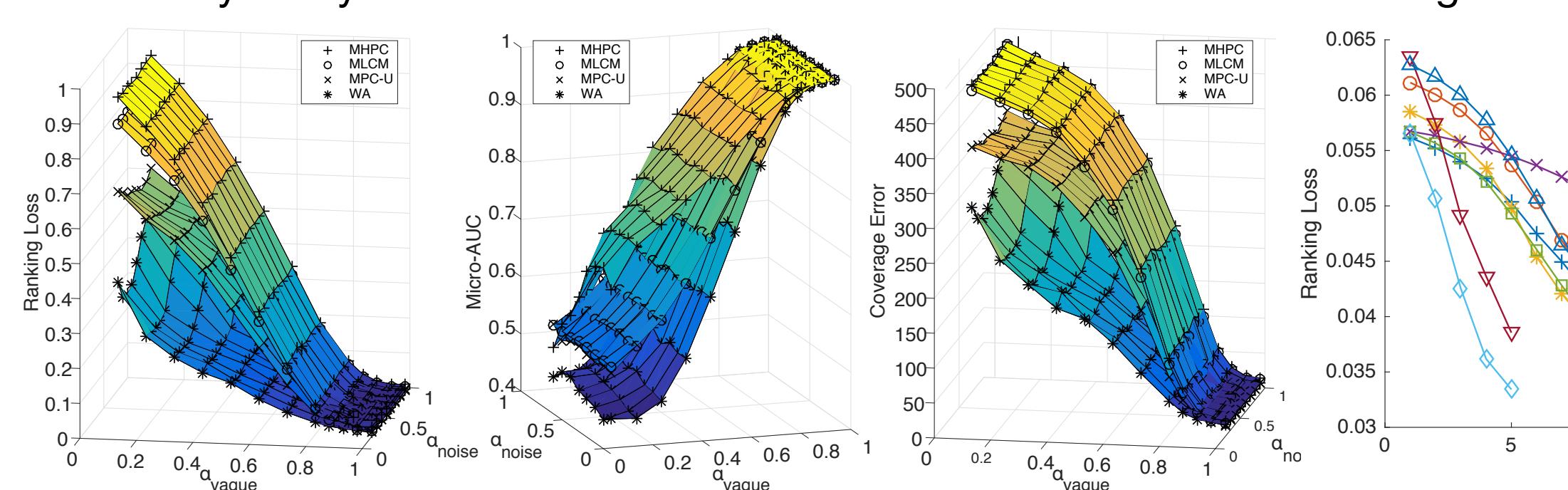
Noisy and ambiguous labels in each annotation:



### Parameter Estimation



### Sensitivity Analysis



### Crowdsourced Online Medical Consultation

Each instance is a set of symptoms that a user describes. For each set of symptoms, disease names are the labels we collected from different doctors.

Instance	Doctor.id	Labels	Ground Truth
sneezing, runny nose,	52****17	{common cold, allergic rhinitis}	{common cold, allergic rhinitis}
sleepy	46****35	{common cold, rhinitis}	{common cold, rhinitis, sinusitis, antritis}
	53****11	{rhinitis, sinusitis}	

Datasets	Methods	Evaluation Metrics		
		Ranking Loss	Micro-AUC	Coverage Error
MED.1	SA	0.520 (5)	0.620 (5)	213.175 (5)
	WA	0.518 (4)	0.621 (3)	212.844 (4)
	MPC-U	0.304 (3)	0.547 (4)	125.108 (3)
	MlCM	0.262 (2)	0.643 (2)	108.146 (1)
MED.2	MHPC	0.196 (1)	0.754 (1)	125.107 (2)
	SA	0.358 (5)	0.603 (4)	75.544 (5)
	WA	0.357 (4)	0.604 (3)	75.416 (4)
	MPC-U	0.200 (2)	0.539 (5)	42.630 (3)
MED.3	MlCM	0.268 (3)	0.680 (2)	31.221 (1)
	MHPC	0.166 (1)	0.715 (1)	35.336 (2)
	SA	0.067 (4)	0.706 (2)	3.400 (4)
	WA	0.065 (3)	0.704 (3)	3.314 (3)
MED.4	MPC-U	0.069 (5)	0.432 (5)	3.543 (5)
	MlCM	0.064 (2)	0.542 (4)	3.288 (2)
	MHPC	0.038 (1)	0.847 (1)	2.000 (1)
	SA	0.301 (5)	0.601 (4)	38.325 (5)
MED.5	WA	0.300 (4)	0.602 (3)	38.275 (4)
	MPC-U	0.173 (2)	0.460 (5)	22.242 (3)
	MlCM	0.225 (3)	0.617 (2)	19.325 (2)
	MHPC	0.118 (1)	0.707 (1)	15.233 (1)
MED.6	SA	0.355 (5)	0.563 (4)	67.534 (5)
	WA	0.353 (4)	0.564 (3)	67.170 (4)
	MPC-U	0.189 (1)	0.549 (5)	36.080 (2)
	MlCM	0.276 (3)	0.565 (2)	52.523 (3)
MED.7	MHPC	0.232 (2)	0.572 (1)	32.648 (1)
	SA	0.365 (5)	0.598 (3)	47.175 (5)
	WA	0.363 (4)	0.599 (2)	46.991 (4)
	MPC-U	0.236 (3)	0.491 (5)	30.632 (3)
MED.8	MlCM	0.190 (1)	0.632 (1)	24.798 (2)
	MHPC	0.203 (2)	0.594 (4)	22.633 (1)

Table 1: Performance on medical data sets.