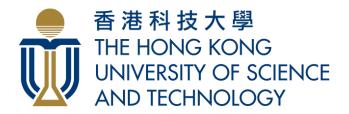
# Relational Deep Learning: A Deep Latent Variable Model for Link Prediction

#### Hao Wang, Xingjian Shi, Dit-Yan Yeung





#### Motivation

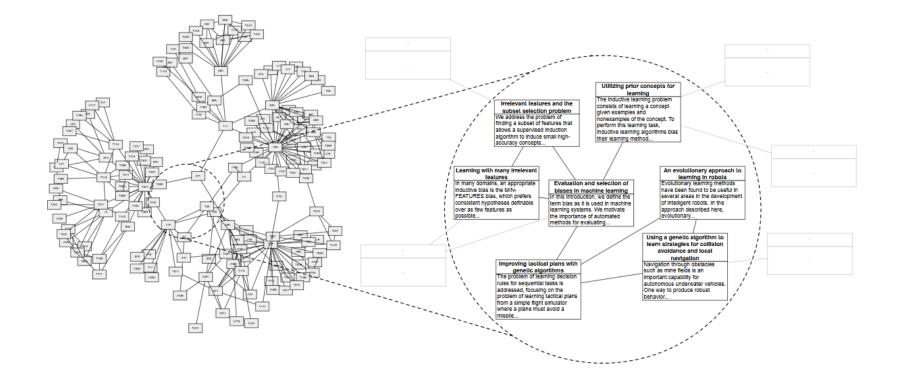
- Bayesian Deep Learning
- •Relational Deep Learning
- Parameter Learning
- •Experiments
- Conclusion

### **Motivation: Link Prediction**



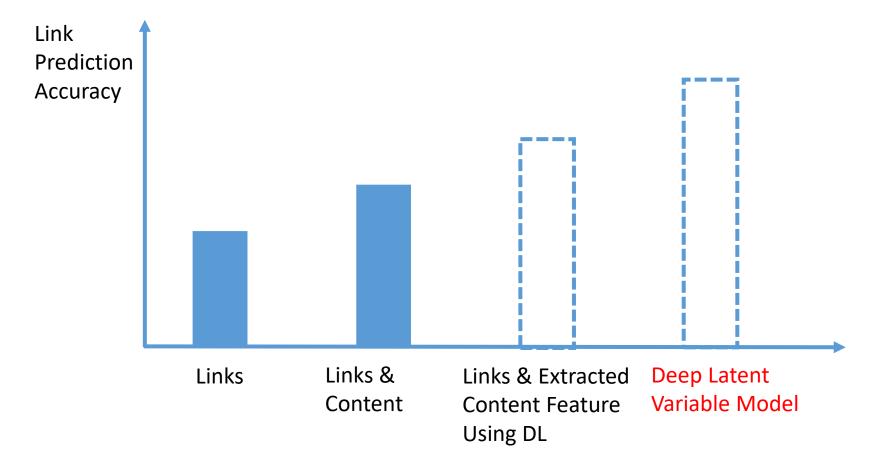
Social Network Analysis (e.g., prediction friendship in Facebook)

### **Motivation: Link Prediction**

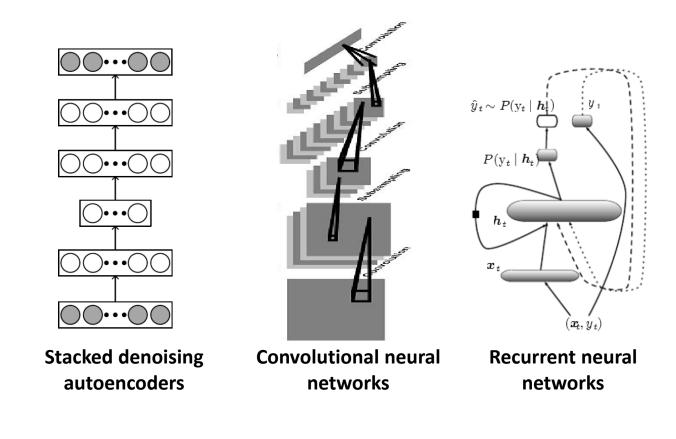


Document Networks (e.g., citation networks, co-author networks)

### **Motivation: Deep Latent Variable Models**



#### **Motivation: Deep Latent Variable Models**

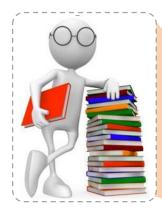


# **Typically for i.i.d. data**

#### Motivation

- Bayesian Deep Learning
- •Relational Deep Learning
- Parameter Learning
- •Experiments
- Conclusion

# **Bayesian Deep Learning**



#### **Perception component**

Content understanding Posts by users Text in articles





#### Task-Specific component

Target task Link prediction

#### **Bayesian deep learning (BDL)**

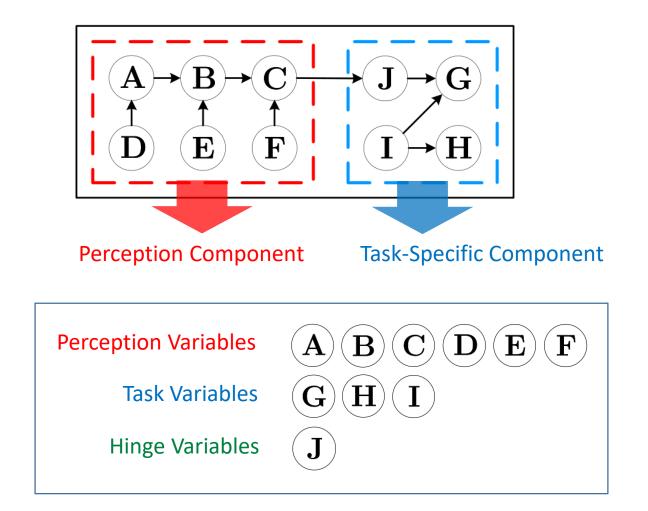
- •Maximum a posteriori (MAP)
- •Markov chain Monte Carlo (MCMC)
- •Variational inference (VI)

[Wang et al. 2016]

### **Bayesian Deep Learning**

| Applications | Models                                   | Hinge Variables                      | Learning              |
|--------------|--|--------------------------------------|-----------------------|
|              | CDL [Wang et al.]                        | $\{\mathbf{V}\}$                     | MAP                   |
| Recommender  | Bayesian CDL [Wang et al.]               | $\{\mathbf{V}\}$                     | Gibbs Sampling        |
| Systems      | Marginalized CDL [Li et al.]             | $\{\mathbf{V}\}$                     | MAP                   |
| Systems      | Symmetric CDL [Li et al.]                | $\{\mathbf{V},\mathbf{U}\}$          | MAP                   |
|              | Collaborative Deep Ranking [Ying et al.] | $\{\mathbf{V}\}$                     | MAP                   |
| Topic        | Relational SDAE [Wang et al.]            | $\{\mathbf{S}\}$                     | MAP                   |
| Models       | DPFA-SBN [Gan et al.]                    | $\{\mathbf{X}\}$                     | Hybrid MC             |
| widdels      | DPFA-RBM [Gan et al.]                    | $\{\mathbf{X}\}$                     | Hybrid MC             |
| Control      | Embed to Control [Watter et al.]         | $\{\mathbf{z}_t, \mathbf{z}_{t+1}\}$ | Variational Inference |

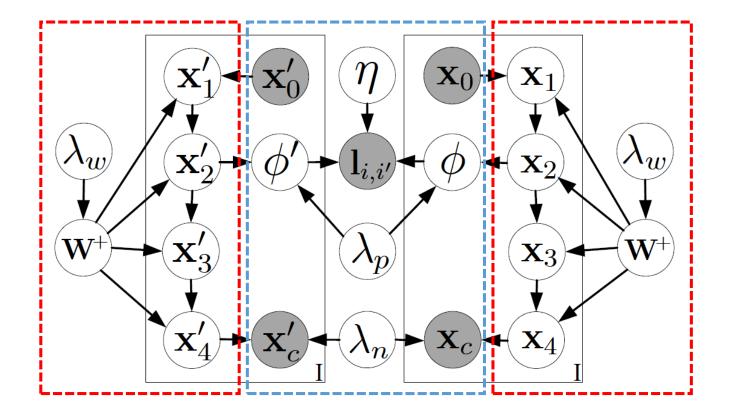
### **A Principled Probabilistic Framework**



[Wang et al. 2016]

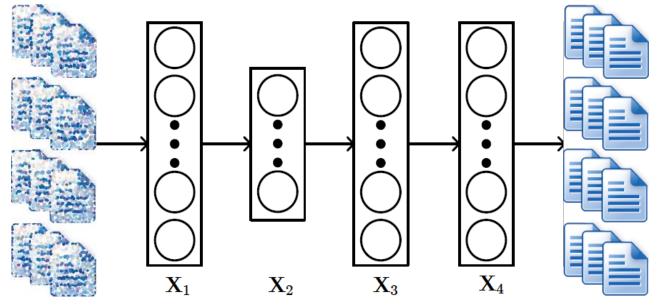
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### **Relational Deep Learning: Graphical Model**



Perception component: relational and deep representation learning Task-specific component: link prediction

# **Stacked Denoising Autoencoders (SDAE)**



Corrupted input

**Clean input** 

SDAE solves the following optimization problem:

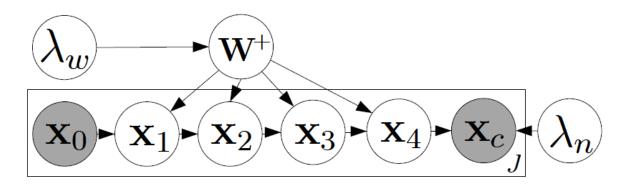
$$\min_{\{\mathbf{W}_l\},\{\mathbf{b}_l\}} \|\mathbf{X}_c - \mathbf{X}_L\|_F^2 + \lambda \sum_l \|\mathbf{W}_l\|_F^2,$$

where  $\lambda$  is a regularization parameter and  $\|\cdot\|_F$  denotes the Frobenius norm.

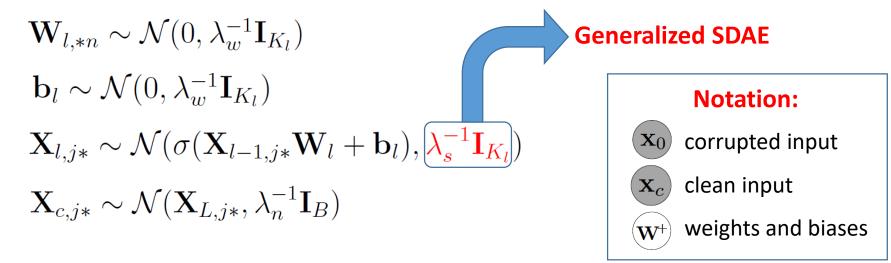
[Vincent et al. 2010]

## **Probabilistic SDAE**

#### **Graphical model:**

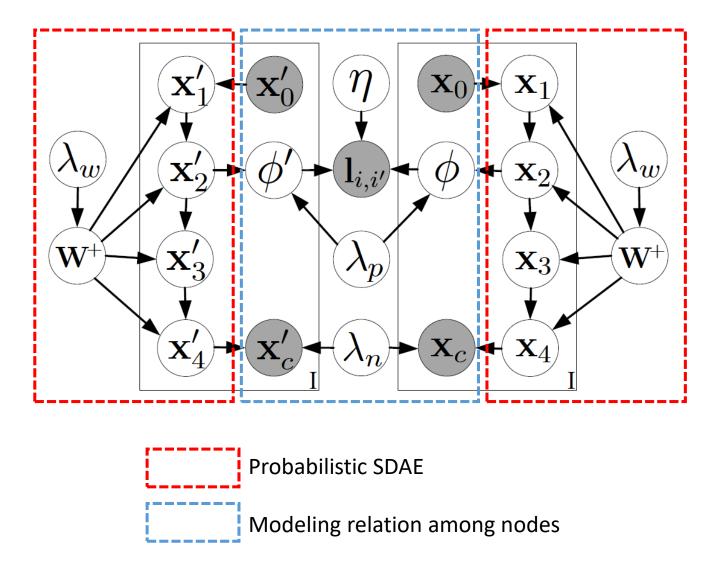


#### **Generative process:**

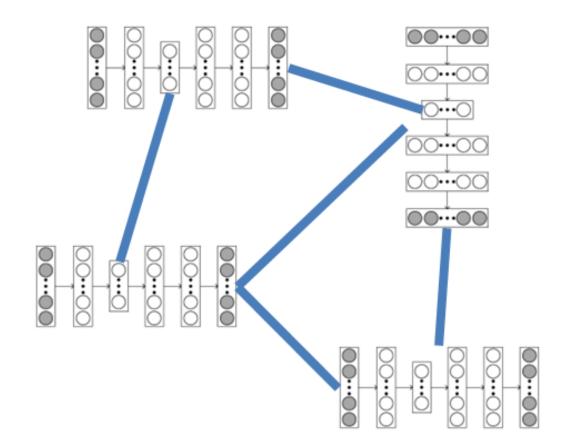


[Wang et al. 2015]

### **Relational Deep Learning**



### **Network of Probabilistic SDAE**



Many interconnected probabilistic SDAEs with shared weights

- Motivation
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maximizing the posterior probability is equivalent to maximizing the joint log-likelihood

$$\begin{aligned} \mathscr{L} &= -\frac{\lambda_w}{2} \sum_l (\|\mathbf{W}_l\|_F^2 + \|\mathbf{b}_l\|_2^2) \\ &- \frac{\lambda_p}{2} \sum_i \|\boldsymbol{\phi}_i - \mathbf{X}_{\frac{L}{2},i*}^T\|_2^2 - \frac{\lambda_n}{2} \sum_i \|\mathbf{X}_{L,i*} - \mathbf{X}_{c,i*}\|_2^2 \\ &- \frac{\lambda_s}{2} \sum_l \sum_i \|\sigma(\mathbf{X}_{l-1,i*}\mathbf{W}_l + \mathbf{b}_l) - \mathbf{X}_{l,i*}\|_2^2 \\ &- \frac{\lambda_e}{2} \|\boldsymbol{\eta}\|_2^2 + \sum_{l_{i,i'}=1} \log \sigma(\boldsymbol{\eta}^T(\boldsymbol{\phi}_i \circ \boldsymbol{\phi}_{i'})). \end{aligned}$$

Prior (regularization) for link prediction parameters, weights, and biases

$$\mathscr{L} = -\frac{\lambda_w}{2} \sum_l (\|\mathbf{W}_l\|_F^2 + \|\mathbf{b}_l\|_2^2)$$
$$-\frac{\lambda_p}{2} \sum_i \|\boldsymbol{\phi}_i - \mathbf{X}_{\frac{L}{2},i*}^T\|_2^2 - \frac{\lambda_n}{2} \sum_i \|\mathbf{X}_{L,i*} - \mathbf{X}_{c,i*}\|_2^2$$
$$-\frac{\lambda_s}{2} \sum_l \sum_i \|\sigma(\mathbf{X}_{l-1,i*}\mathbf{W}_l + \mathbf{b}_l) - \mathbf{X}_{l,i*}\|_2^2$$
$$-\frac{\lambda_e}{2} \|\boldsymbol{\eta}\|_2^2 + \sum_{l_{i,i'}=1} \log \sigma(\boldsymbol{\eta}^T(\boldsymbol{\phi}_i \circ \boldsymbol{\phi}_{i'})).$$

Generating node features from content representation with Gaussian offset

$$\mathcal{L} = -\frac{\lambda_w}{2} \sum_l (\|\mathbf{W}_l\|_F^2 + \|\mathbf{b}_l\|_2^2)$$
$$-\frac{\lambda_p}{2} \sum_i \|\boldsymbol{\phi}_i - \mathbf{X}_{\frac{L}{2},i*}^T\|_2^2 - \frac{\lambda_n}{2} \sum_i \|\mathbf{X}_{L,i*} - \mathbf{X}_{c,i*}\|_2^2$$
$$-\frac{\lambda_s}{2} \sum_l \sum_i \|\sigma(\mathbf{X}_{l-1,i*}\mathbf{W}_l + \mathbf{b}_l) - \mathbf{X}_{l,i*}\|_2^2$$
$$-\frac{\lambda_e}{2} \|\boldsymbol{\eta}\|_2^2 + \sum_{l_{i,i'}=1} \log \sigma(\boldsymbol{\eta}^T(\boldsymbol{\phi}_i \circ \boldsymbol{\phi}_{i'})).$$

'Generating' clean input from the output of probabilistic SDAE with Gaussian offset

$$\begin{aligned} \mathscr{L} &= -\frac{\lambda_w}{2} \sum_l (\|\mathbf{W}_l\|_F^2 + \|\mathbf{b}_l\|_2^2) \\ &- \frac{\lambda_p}{2} \sum_i \|\boldsymbol{\phi}_i - \mathbf{X}_{\frac{L}{2},i*}^T\|_2^2 - \frac{\lambda_n}{2} \sum_i \|\mathbf{X}_{L,i*} - \mathbf{X}_{c,i*}\|_2^2 \\ &- \frac{\lambda_s}{2} \sum_l \sum_i \|\sigma(\mathbf{X}_{l-1,i*}\mathbf{W}_l + \mathbf{b}_l) - \mathbf{X}_{l,i*}\|_2^2 \\ &- \frac{\lambda_e}{2} \|\boldsymbol{\eta}\|_2^2 + \sum_{l_{i,i'}=1} \log \sigma(\boldsymbol{\eta}^T(\boldsymbol{\phi}_i \circ \boldsymbol{\phi}_{i'})). \end{aligned}$$

Generating the input of Layer I from the output of Layer I-1 with Gaussian offset

$$\begin{aligned} \mathscr{L} &= -\frac{\lambda_w}{2} \sum_l (\|\mathbf{W}_l\|_F^2 + \|\mathbf{b}_l\|_2^2) \\ &- \frac{\lambda_p}{2} \sum_i \|\boldsymbol{\phi}_i - \mathbf{X}_{\frac{L}{2},i*}^T\|_2^2 - \frac{\lambda_n}{2} \sum_i \|\mathbf{X}_{L,i*} - \mathbf{X}_{c,i*}\|_2^2 \\ &- \frac{\lambda_s}{2} \sum_l \sum_i \|\sigma(\mathbf{X}_{l-1,i*}\mathbf{W}_l + \mathbf{b}_l) - \mathbf{X}_{l,i*}\|_2^2 \\ &- \frac{\lambda_e}{2} \|\boldsymbol{\eta}\|_2^2 + \sum_{l_{i,i'}=1} \log \sigma(\boldsymbol{\eta}^T(\boldsymbol{\phi}_i \circ \boldsymbol{\phi}_{i'})). \end{aligned}$$

Generating links from Bernoulli distributions parameterized by  $\eta$  and  $\varphi$ 

$$\begin{aligned} \mathscr{L} &= -\frac{\lambda_w}{2} \sum_l (\|\mathbf{W}_l\|_F^2 + \|\mathbf{b}_l\|_2^2) \\ &- \frac{\lambda_p}{2} \sum_i \|\boldsymbol{\phi}_i - \mathbf{X}_{\frac{L}{2},i*}^T\|_2^2 - \frac{\lambda_n}{2} \sum_i \|\mathbf{X}_{L,i*} - \mathbf{X}_{c,i*}\|_2^2 \\ &- \frac{\lambda_s}{2} \sum_l \sum_i \|\sigma(\mathbf{X}_{l-1,i*}\mathbf{W}_l + \mathbf{b}_l) - \mathbf{X}_{l,i*}\|_2^2 \\ &- \frac{\lambda_e}{2} \|\boldsymbol{\eta}\|_2^2 + \sum_{l_{i,i'}=1} \log \sigma(\boldsymbol{\eta}^T(\boldsymbol{\phi}_i \circ \boldsymbol{\phi}_{i'})) \end{aligned}$$

# **Bayesian Treatment: Generalized Variational Inference**

 $\log q_j^*(\mathbf{Z}_j) = \mathcal{E}_{i \neq j}[\log p(\mathbf{X}_0, \mathbf{X}_c, \mathbf{Z})] + const$ 

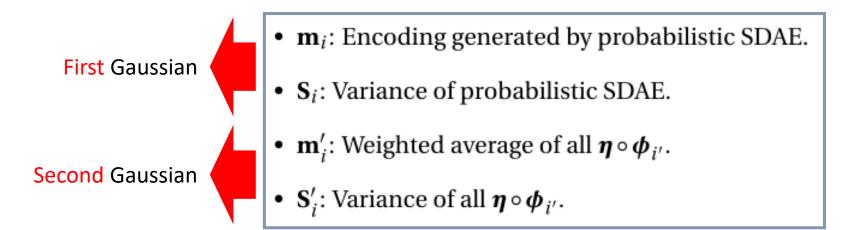
- $q_1(\mathbf{Z}_1) = q(\mathbf{W}^+)$ : Variational distributions for weights/biases.
- $q_2(\mathbf{Z}_2) = q(\boldsymbol{\phi}_i)$ : Variational distributions for generated node features.
- $q_3(\mathbf{Z}_3) = q(\mathbf{\eta})$ : Variational distributions for parameters of the link prediction model.
- $q_4(\mathbf{Z}_4) = q(\xi)$ : Variational parameters to approximate the sigmoid function.

Use Laplace approximation rather than variational inference for weights/biases.

#### Example: Updating $\phi$ as a Product of Gaussians

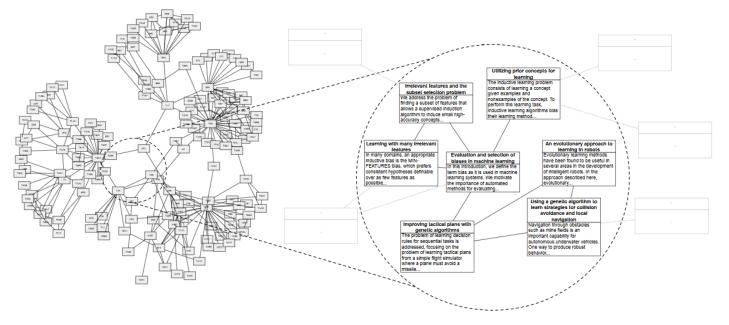
Update  $\varphi$  for node i as a product of two Gaussians

$$q(\boldsymbol{\phi}_i | \mathbf{X}_{0,i*}) \approx \mathcal{N}(\boldsymbol{\phi}_i | \boldsymbol{\mu}_i, \boldsymbol{\Sigma}_i)$$
$$\boldsymbol{\mu}_i = \boldsymbol{\Sigma}_i (\mathbf{S}_i^{-1} \mathbf{m}_i + {\mathbf{S}'_i}^{-1} \mathbf{m}'_i)$$
$$\boldsymbol{\Sigma}_i^{-1} = \mathbf{S}_i^{-1} + {\mathbf{S}'_i}^{-1}.$$



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#### **Experiments: Settings**



#### Document Networks (e.g., citation networks)

| datasets    | # nodes | # links |  |  |
|-------------|---------|---------|--|--|
| citeulike-a | 16,980  | 44,709  |  |  |
| citeulike-t | 25,975  | 32,565  |  |  |
| arXiv       | 27,770  | 352,807 |  |  |

#### **Experiments: Link Rank and AUC**

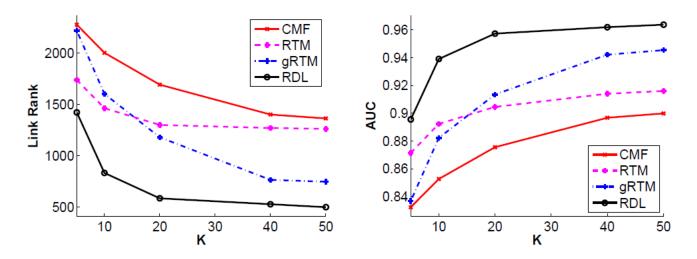


Figure 2: Link rank and AUC of compared models for *citeulike-a*. A 2-layer RDL is used.

Link rank: how high our predicted links rank in the ground truth AUC: area under curve

#### **Experiments: Link Rank and AUC**

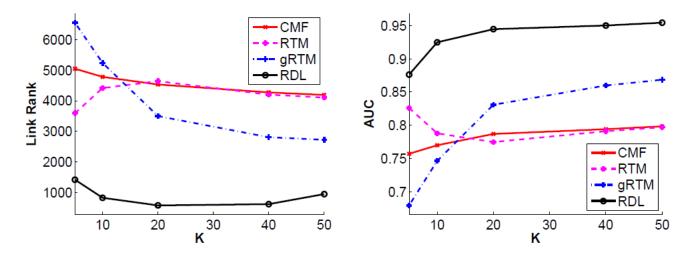


Figure 3: Link rank and AUC of compared models for *citeulike-t*. A 2-layer RDL is used.

Link rank: how high our predicted links rank in the ground truth AUC: area under curve

### **Experiments: Link Rank and AUC**

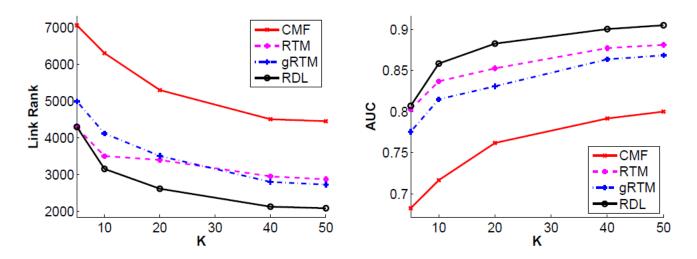


Figure 4: Link rank and AUC of compared models for *arXiv*. A 2-layer RDL is used.

Link rank: how high our predicted links rank in the ground truth AUC: area under curve

### **Experiments: RDL Variants**

#### Link rank of baselines (the first 3 columns) and RDL variants (the last 4 columns) on three datasets (L = 4)

| Method      | VAE+BLR | VFAE+BLR | SDAE+BLR | MAPRDL  | BSDAE1+BLR | BSDAE2+BLR | BayesRDL |
|-------------|---------|----------|----------|---------|------------|------------|----------|
| citeulike-a | 980.81  | 960.15   | 992.48   | 495.97  | 849.02     | 761.57     | 473.59   |
| citeulike-t | 1599.62 | 1531.16  | 1356.85  | 951.31  | 1341.15    | 1310.12    | 911.31   |
| arXiv       | 3367.25 | 3316.29  | 2916.18  | 2028.72 | 2947.79    | 2708.17    | 1982.84  |

VAE: Variational Autoencoder

VRAE: Variational Fair Autoencoder

**BLR: Bayesian Logistic Regression** 

BSDAE1: Bayesian treatment of probabilistic SDAE (mean only)

BSDAE2: Bayesian treatment of probabilistic SDAE (mean and variance)

MAPRDL: RDL with MAP inference

BayesRDL: RDL with full Bayesian treatment

#### **Experiments: Depth**

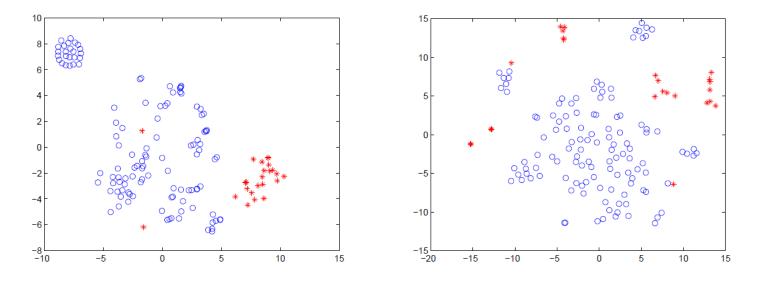
|             | Link Rank |         | AUC     |       |       |       |
|-------------|-----------|---------|---------|-------|-------|-------|
|             | RDL-1     | RDL-2   | RDL-3   | RDL-1 | RDL-2 | RDL-3 |
| citeulike-a | 825.74    | 495.97  | 488.41  | 0.939 | 0.964 | 0.963 |
| citeulike-t | 2060.17   | 951.31  | 912.43  | 0.894 | 0.954 | 0.955 |
| arXiv       | 5241.97   | 2080.72 | 2730.08 | 0.755 | 0.905 | 0.855 |

#### Performance of RDL with different number of layers (MAP)

#### Performance of RDL with different number of layers (Bayesian treatment)

|             | Link Rank |         |         | AUC   |       |       |
|-------------|-----------|---------|---------|-------|-------|-------|
|             | RDL-1     | RDL-2   | RDL-3   | RDL-1 | RDL-2 | RDL-3 |
| citeulike-a | 789.85    | 473.59  | 471.47  | 0.946 | 0.971 | 0.970 |
| citeulike-t | 1904.83   | 911.31  | 867.78  | 0.906 | 0.956 | 0.960 |
| arXiv       | 4965.01   | 1982.84 | 2612.12 | 0.801 | 0.914 | 0.866 |

### **Case Study: RDL and RTM**



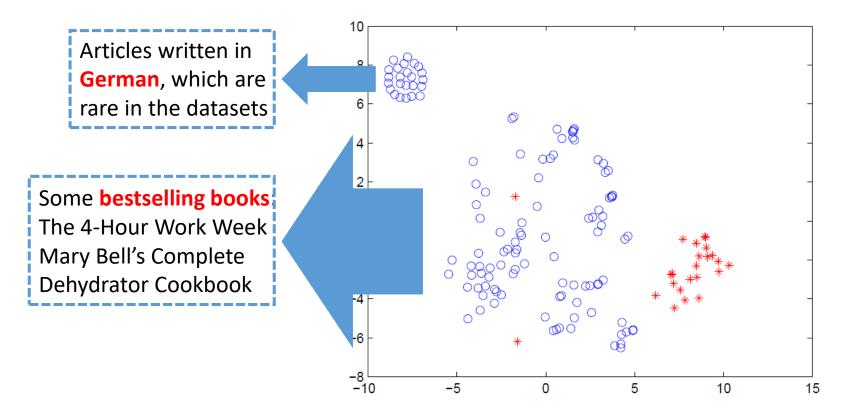
t-SNE visualization of latent factors learned by RDL (left) and RTM (right).

Target article:

From DNA sequence to transcriptional behaviour: a quantitative approach

\* (red): articles with links to the target article
O (blue): articles without links to the target article

## **Case Study: RDL**



t-SNE visualization of latent factors learned by RDL.

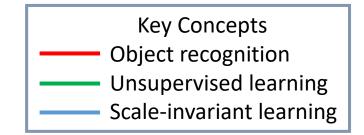
Target article:

From DNA sequence to transcriptional behaviour: a quantitative approach

## Case Study: RDL ang gRTM

#### Top 10 link predictions made by gRTM and RDL for two articles from citeulike-a

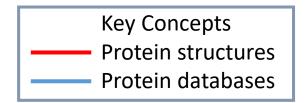
|      | Query: Object class recognition by unsupervised scale-invariant learning   |
|------|--|
|      | Layered depth images   |
|      | Using spin images for efficient object recognition in cluttered 3D scenes  |
|      | Snakes: active contour models  |
|      | Visual learning and recognition of 3-D objects from appearance             |
| m    | Contextual priming for object detection                                    |
| R    | Visual categorization with bags of keypoints                               |
| gRTM | Non-parametric model for background subtraction                            |
| -    | Alignment by maximization of mutual information                            |
|      | Rapid object detection using a boosted cascade of simple features          |
|      | W4: real-time surveillance of people and their activities                  |
|      | Distinctive image features from scale-invariant keypoints                  |
|      | visual learning and recognition of 3-D objects from appearance             |
|      | Object recognition with features inspired by visual cortex                 |
|      | Unsupervised learning of models for recognition                            |
|      | Robust object recognition with cortex-like mechanisms                      |
| RDL  | Generative versus discriminative methods for object recognition            |
| ΗĔ   | Using spin images for efficient object recognition in cluttered 3D scenes  |
|      | Learning generative visual models from few training examples               |
|      | 3D object modeling and recognition using affine-invariant patches          |
|      | A Bayesian approach to unsupervised one-shot learning of object categories |



## Case Study: RDL ang gRTM

#### Top 10 link predictions made by gRTM and RDL for two articles from citeulike-a

|      | Query: SCOP database in 2004: refinements integrate structure and sequence family data     |
|------|--|
|      | Pfam: multiple sequence alignments and HMM-profiles of protein domains                     |
|      | Structure, function and evolution of multidomain proteins                                  |
|      | Greengenes, a chimera-checked 16S rRNA gene database and workbench compatible with ARB     |
|      | Nature of the protein universe   |
| 000  | The CATH domain structure database and related resources                                   |
| , R  | Phylogenetic classification of short environmental DNA fragments                           |
| gRTM | The catalytic site atlas: a resource of catalytic sites and residues identified in enzymes |
| 5    | LGA: a method for finding 3D similarities in protein structures                            |
|      | Amino acid substitution matrices from protein blocks                                       |
|      | Multiple protein sequence alignment  |
|      | The universal protein resource (UniProt)   |
|      | E-MSD: an integrated data resource for bioinformatics                                      |
|      | Gene3D: comprehensive structural and functional annotation of genomes                      |
|      | The universal protein resource (UniProt) in 2010   |
|      | Gene3D: modelling protein structure, function and evolution                                |
| RDL  | The universal protein resource (UniProt): an expanding universe of protein information     |
| Ĕ    | Pfam: clans, web tools and services  |
|      | The Pfam protein families database   |
|      | The protein data bank  |
|      | SCOP: a structural classification of proteins database                                     |



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#### Conclusion

- •First Bayesian DL model for link prediction
- •Joint Bayesian DL is beneficial
- •Significant improvement on the state of the art
- •RDL as representation learning

#### **Future Work**

#### Multi-relational data (co-author & citation networks)

- •Boost predictive performance
- •Discover relationship between different networks

#### •GVI for other neural nets (CNN/RNN) and BayesNets

- •pSDAE + link prediction
- pCNN + recommendation
- •pRNN + community detection

#### •Replace probabilistic SDAE with other Bayesian neural nets

- •Variational autoencoders
- •Natural-parameter networks



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