

Online Collective Inference



Jay Pujara

U. Maryland, College Park

Ben London

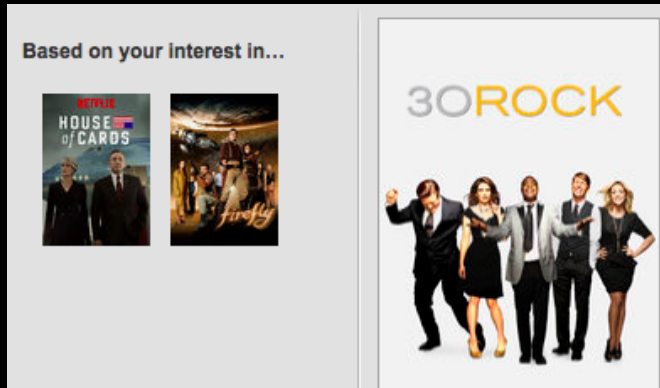
U. Maryland, College Park

Lise Getoor

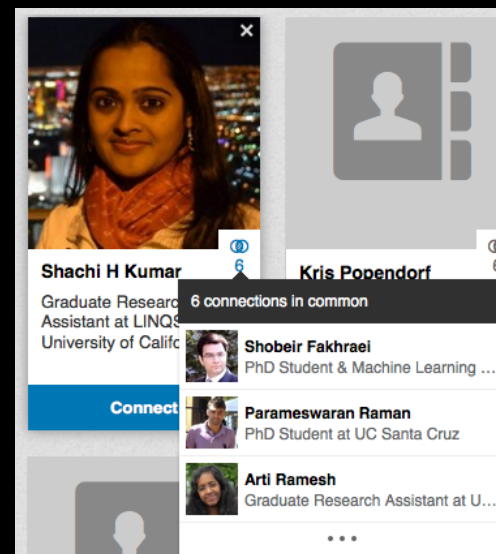
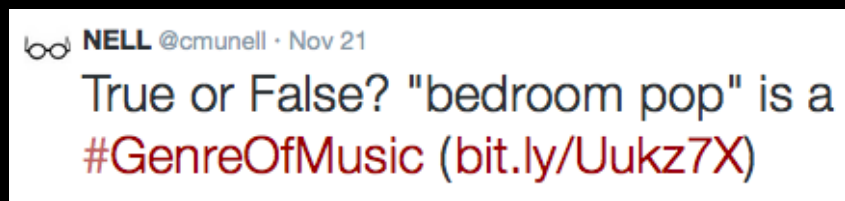
U. California, Santa Cruz



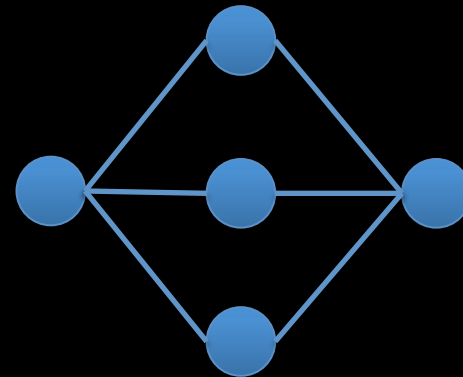
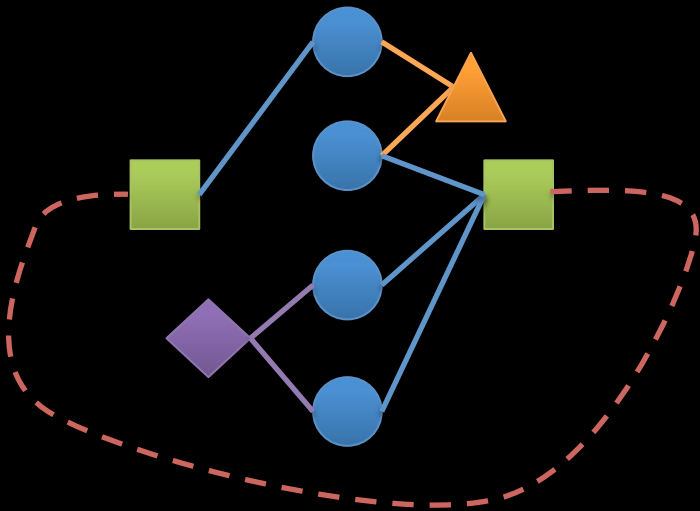
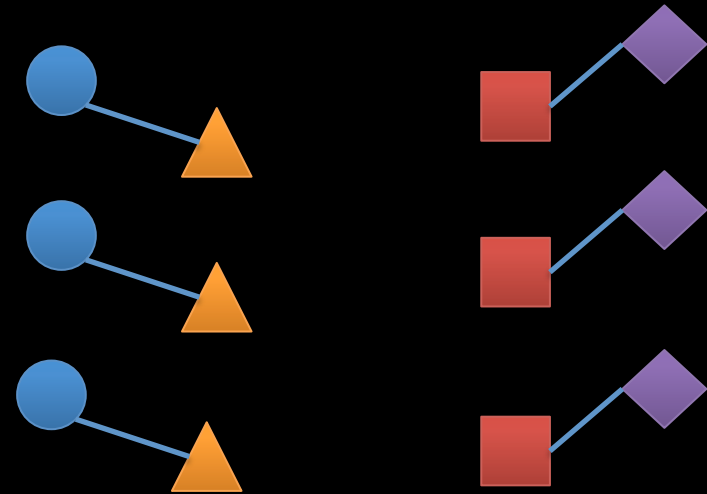
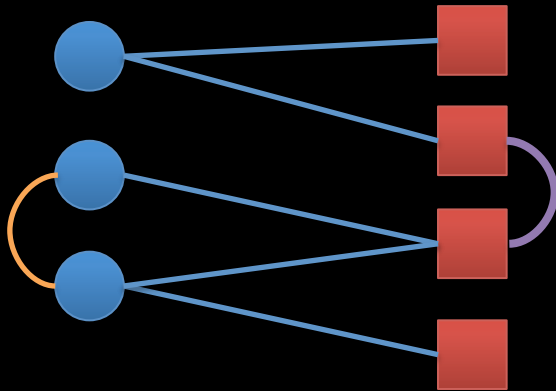
Real-world problems...



Genotype	Genetic Result
CC	Slightly higher odds of disliking the taste of cilantro.
CT	Typical odds of disliking the taste of cilantro.
TT	Slightly lower odds of disliking the taste of cilantro.

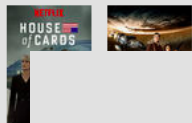


...benefit from relational models



Collaborative Filtering

Based on your interest in...



30ROCK

Based on your interest in...



30ROCK



→ $\text{LIKES}(U1, M2)$



Cow
Cow
→ C

The image shows a screenshot of a LinkedIn interface. On the left, the profile of Shachi H Kumar is visible, with a photo of a woman and a blue 'Connect' button. On the right, the profile of Kris Popendorf is visible, with a grey placeholder for a photo. A central overlay shows '6 connections in common' between them. Below this, a list of shared connections is displayed, including Shobeir Fakhraei, Parameswaran Raman, and Arti Ramesh. To the right of this list, a smaller version of the same interface is shown, with a blue circle and an orange dashed line pointing to it from the right edge of the image.

Shachi H Kumar
Graduate Research Assistant at LINQS
University of California

Kris Popendorf

6 connections in common

- Shobeir Fakhraei**
PhD Student & Machine Learning ...
- Parameswaran Raman**
PhD Student at UC Santa Cruz
- Arti Ramesh**
Graduate Research Assistant at U...

Connect

Knowledge Graph Identification

NELL @cmunell · Nov 21

True or False? "bedroom pop" is a
#GenreOfMusic (bit.ly/Uukz7X)

NELL @cmunell · Nov 21

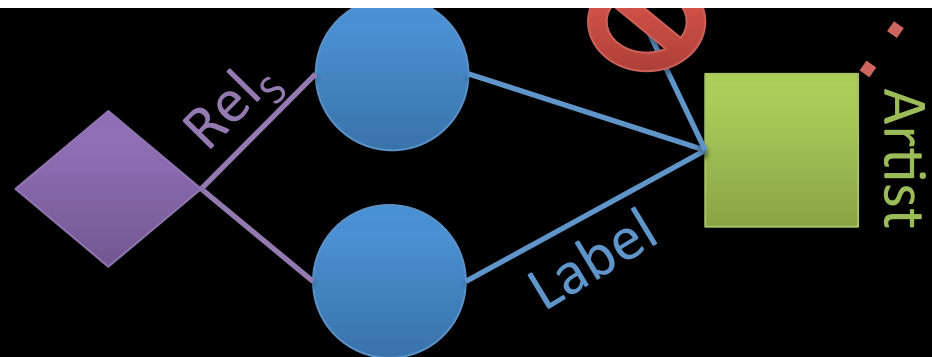
True or False? "bedroom pop" is a
#GenreOfMusic (bit.ly/Uukz7X)

→ LABEL(E2, L)

MUTEX(L1, L2) ∧

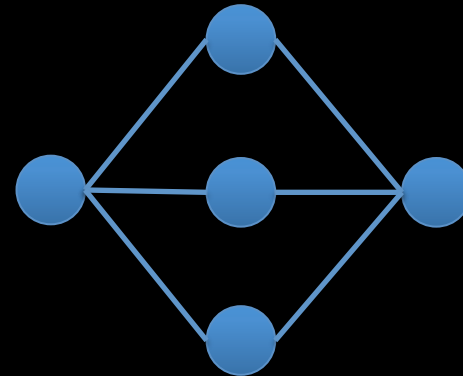
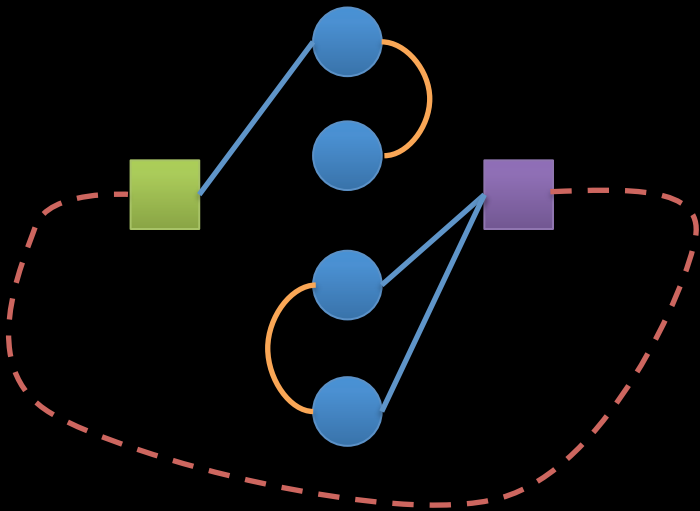
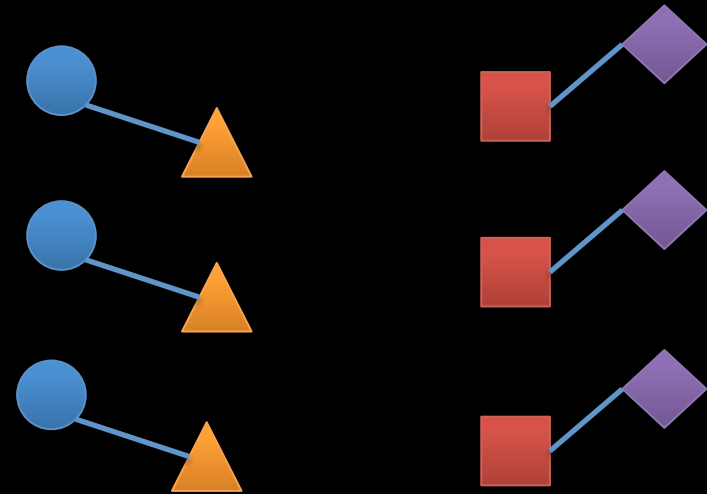
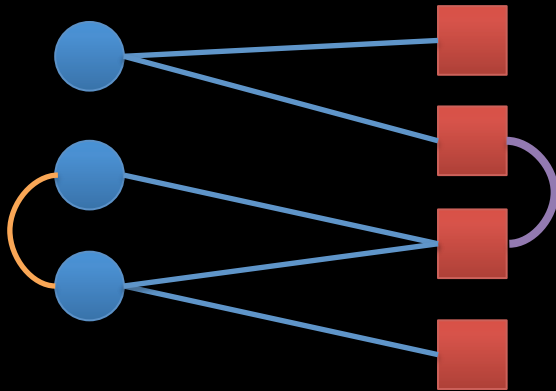
LABEL(E, L1) ∧

→ ¬LABEL(E, L2)



(Jiang et al., ICDM12; Pujara et al., ISWC13)

...benefit from relational models



Real-world problems are big!

Millions of users,
thousands movies

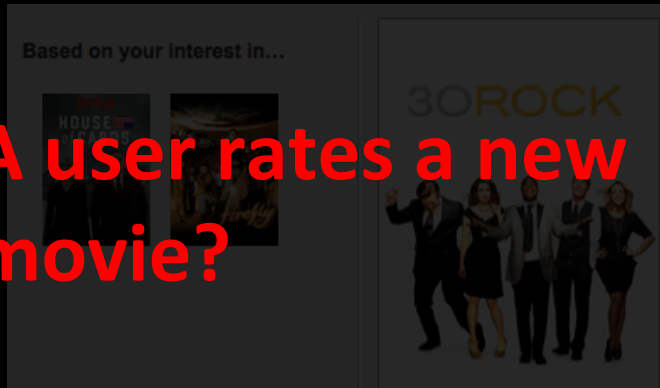
Millions of users,
thousands of genes

Millions of facts,
thousands of
ontological constraints

Millions of users

What happens when?

A user rates a new movie?



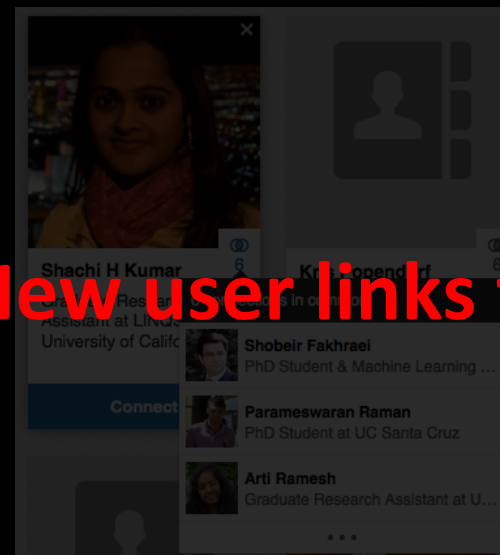
A new genetic similarity is discovered?

Genotype	Genetic Result
CC	Slightly higher odds of disliking the taste of cilantro.
TT	Slightly lower odds of disliking the taste of cilantro.

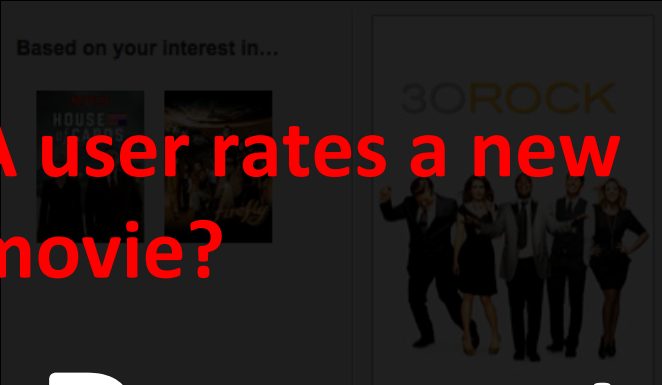
New facts are extracted from the Web?



New user links form?



What happens when?



Based on your interest in...

A user rates a new movie?

The image shows a movie recommendation interface. On the left, there are two movie posters: 'The House' and 'The Fast and the Furious'. On the right, there is a larger poster for '30 Rock'.



A new genetic similarity is discovered?

Genotype	Genetic Result
CC	Slightly higher odds of disliking the taste of cilantro.
CT	Typical odds of disliking the taste of cilantro.
TT	Slightly lower odds of disliking the taste of cilantro.

The image shows a table with two columns: 'Genotype' and 'Genetic Result'. The rows show different genotypes (CC, CT, TT) and their corresponding results regarding the odds of disliking the taste of cilantro.

Repeat Inference!



New facts are extracted from the Web?

The image shows a social media post with the text: 'True or False? "bedroom non" is a #GenreOfMusic (bit.ly/Uukz7X)'.



New user links form?

The image shows a user profile page with a list of users to connect to. The users listed are Shachi H Kumar, Shobeir Fakhraei, Parameswaran Raman, and Arti Ramesh.

Why can't we repeat inference?

- We want rich, collective models!
- But, 10M-1B factors = 1-100s hours*
- Ideal: Inference time balances update cycle
- Insanity is doing the same thing over and over...

Online Collective Inference

PROBLEM SETTING

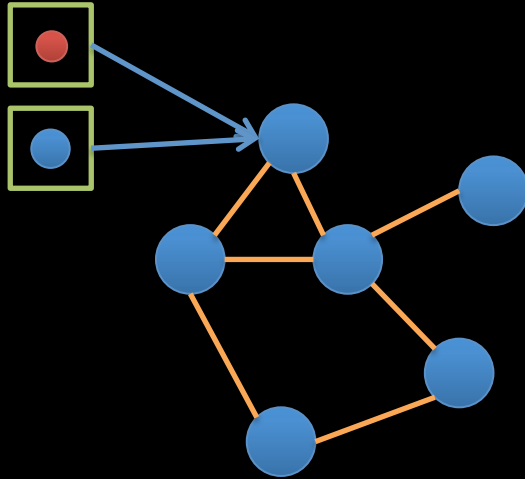
Key Problem

- Real-world problems -> large graphical models
- Changing evidence -> repeat inference

Key Problem

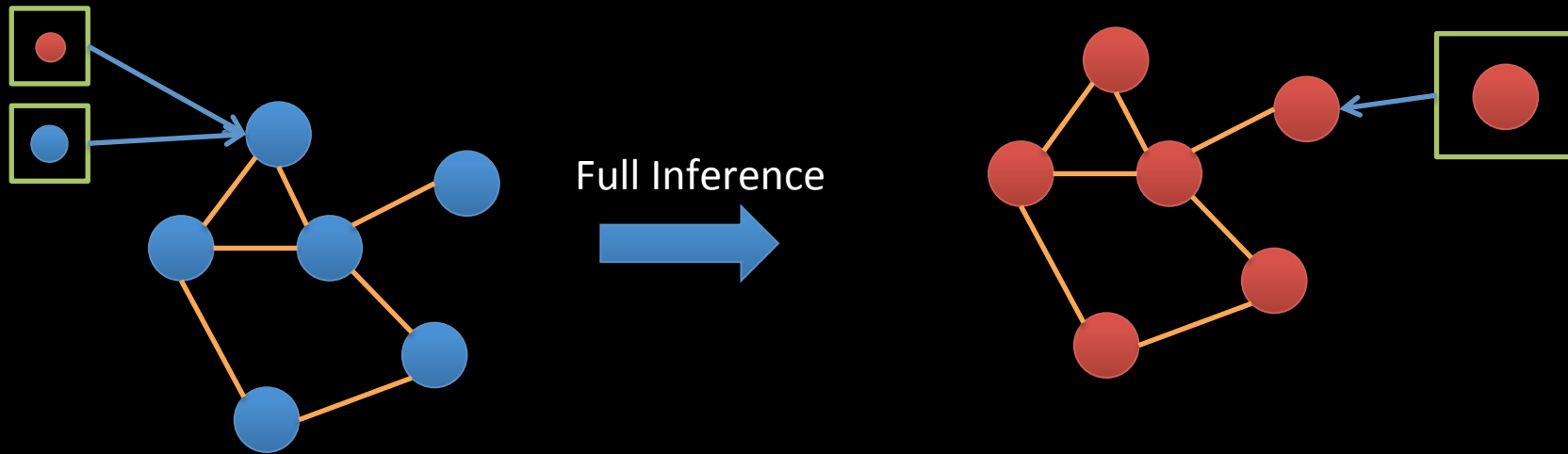
- Real-world problems -> large graphical models
- Changing evidence -> ~~repeat inference~~
- What happens when partially updating inference?
- Can we scalably approximate the MAP state without recomputing inference?

Generic Answer: NO!



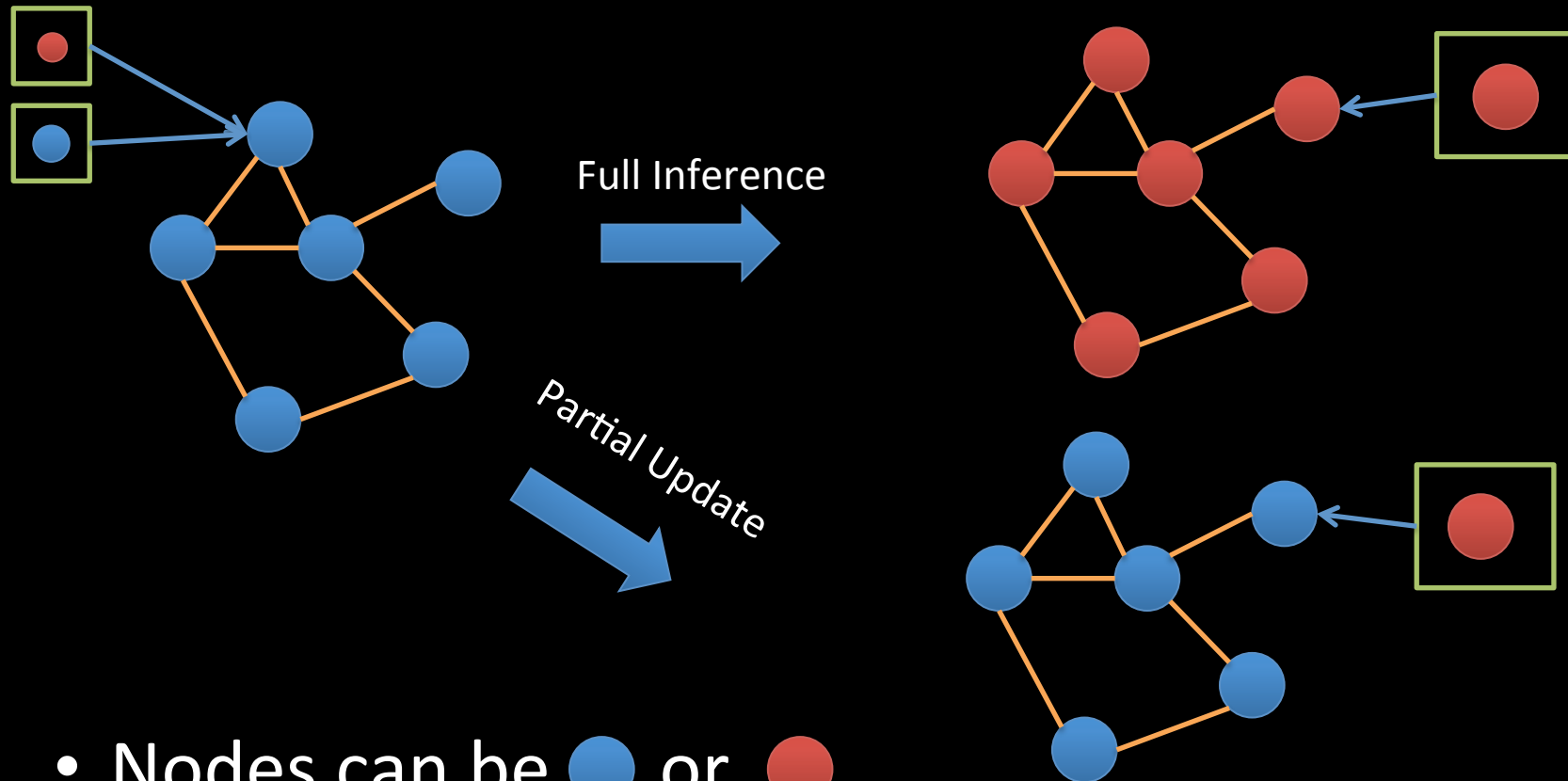
- Nodes can be ● or ●
- Model has prob. mass only when nodes same
- Fix some nodes to ● then observe evidence for ●

Generic Answer: NO!



- Nodes can be ● or ●
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Generic Answer: NO!



- Nodes can be ● or ●
- Model has prob. mass only when nodes same
- Fix some nodes to ● then observe evidence for ●

Previous Work

- Belief Revision
 - e.g. Gardenfors, 1992
- Bayesian Network Updates
 - e.g. Buntine, 1991; Friedman & Goldszmidt, 1997
- Dynamic / Sequential Models
 - e.g. Murphy, 2002 / Fine et al., 1998
- Adaptive Inference
 - e.g. Acar et al., 2008
- BP Message Passing
 - e.g. Nath & Domingos, 2010
- Collective Stability
 - e.g. London et al., 2013

Problem Setting

- Fixed model: dependencies & weights known
- Online: changing evidence or observations
- Closed world: all variables identified
- Budget: infer only m variables in each epoch
- **Strongly-convex inference objective** (e.g. PSL)

Questions:

- What guarantees can we offer?
- Which m variables should we infer?

Approach

- Define “regret” for online collective inference
- Introduce regret bounds for strongly convex inference objectives (like PSL!)
- Develop algorithms to *activate* a subset of the variables during inference, given budget

Online Collective Inference

REGRET BOUNDS

Inference Regret

- General inference problem: estimate $P(Y|X)$
- In online collective inference: fix Y_S , infer $Y_{\overline{S}}$
- Regret (learning): captures distance to optimal
- Regret (inference): the distance between the full inference result and the partial inference update (when conditioning on Y_S)

Defining Regret

- **Regret**: distance between **full** & **approximate** inference

$$\mathcal{R}_n(\mathbf{x}, \mathbf{y}_{\mathcal{S}}; \dot{\mathbf{w}}) \triangleq \frac{1}{n} \| \mathbf{h}(\mathbf{x}; \dot{\mathbf{w}}) - \mathbf{h}(\mathbf{x}, \mathbf{y}_{\mathcal{S}}; \dot{\mathbf{w}}) \|_1$$

where

$$h(\mathbf{x}; \dot{\mathbf{w}}) = \arg \min_{\mathbf{y}} \mathbf{w} \cdot \mathbf{f}(\mathbf{x}, \mathbf{y}) + \frac{\overset{\text{Prior weight}}{w_p}}{2} \|\mathbf{y}\|_2^2.$$

Regret Bound

$$\mathcal{R}_n(\mathbf{x}, \mathbf{y}_{\mathcal{S}}; \dot{\mathbf{w}}) \leq O \left(\sqrt{\frac{B \|\mathbf{w}\|_2}{n \cdot w_p} \|\mathbf{y}_{\mathcal{S}} - \hat{\mathbf{y}}_{\mathcal{S}}\|_1} \right)$$

Regret Ingredients:

Lipschitz constant

2-norm of model weights

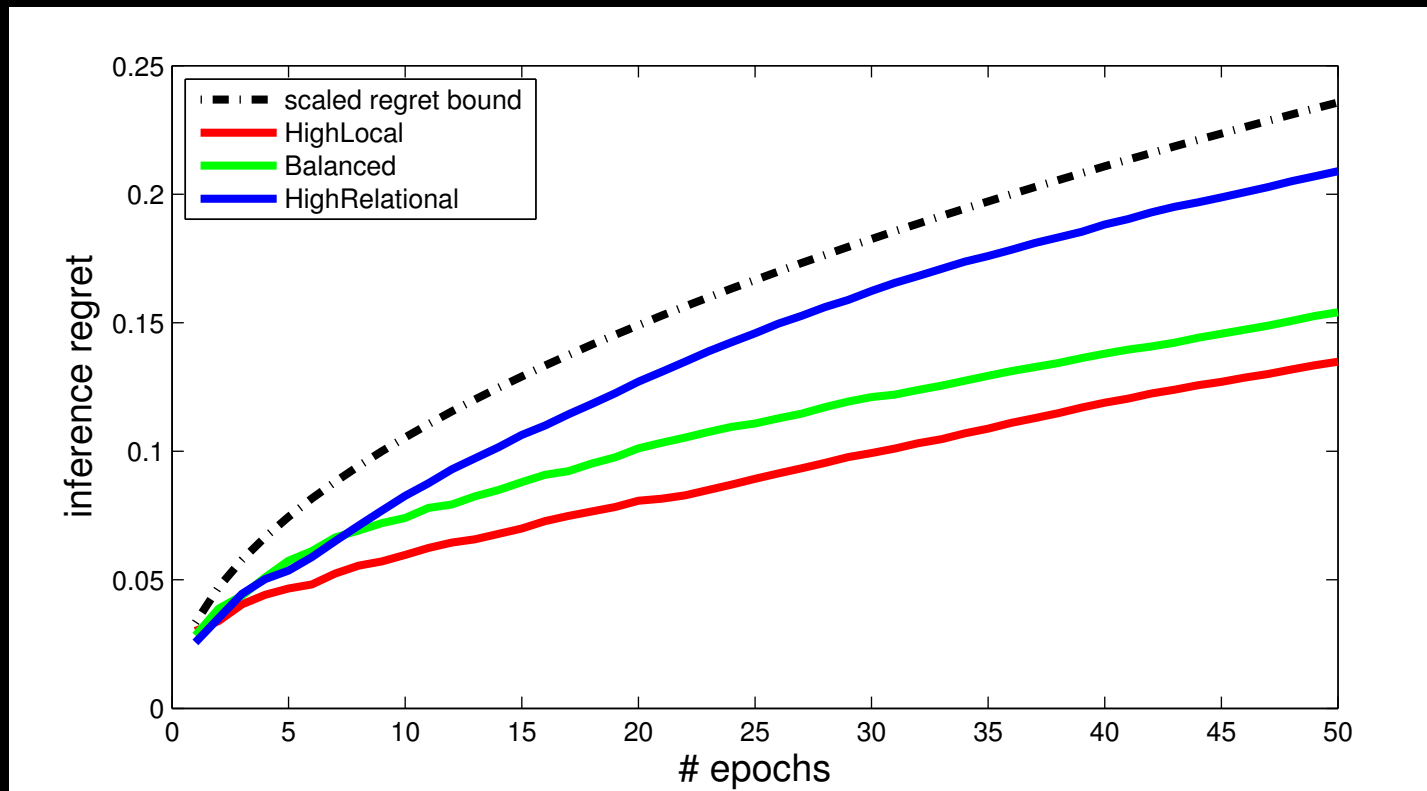
Weight of L_2 prior

L_1 distance fixed variables and
values in full inference

Key Takeaway:

Regret depends on L_1
distance between
fixed variables &
their “true” values in
the MAP state

Validating Regret Bounds



Measure regret of no updates versus full inference, varying the importance of relational features

Online Collective Inference

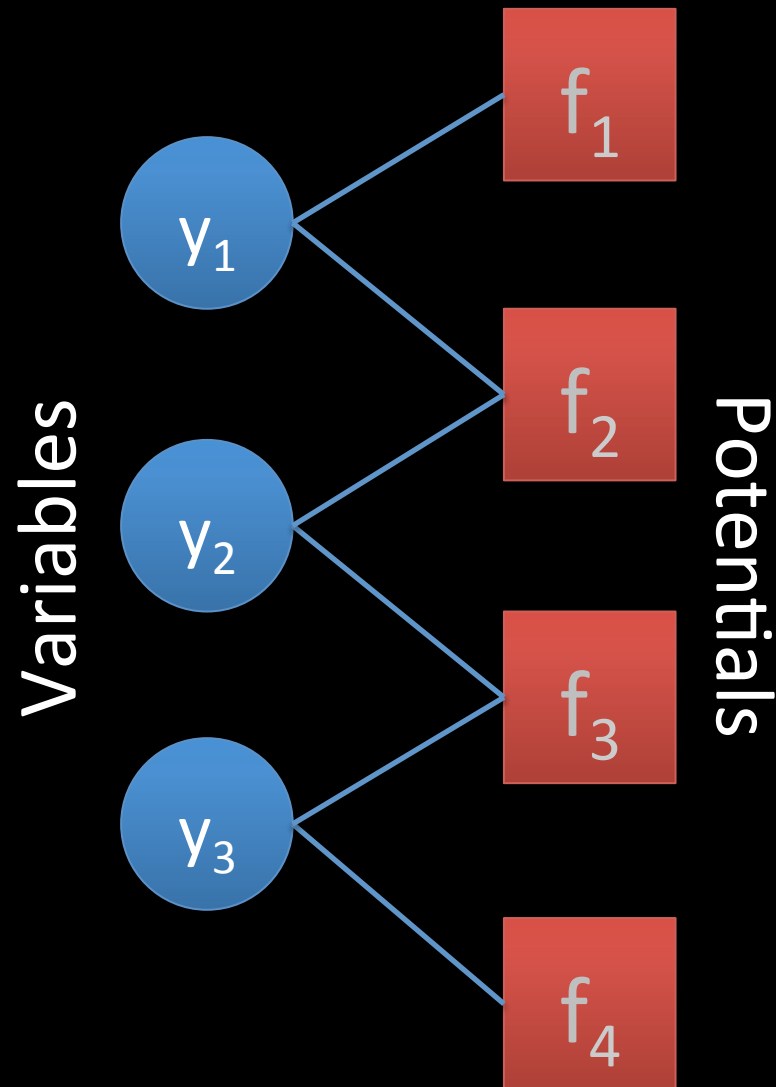
ACTIVATION ALGORITHMS

Which variables to fix?

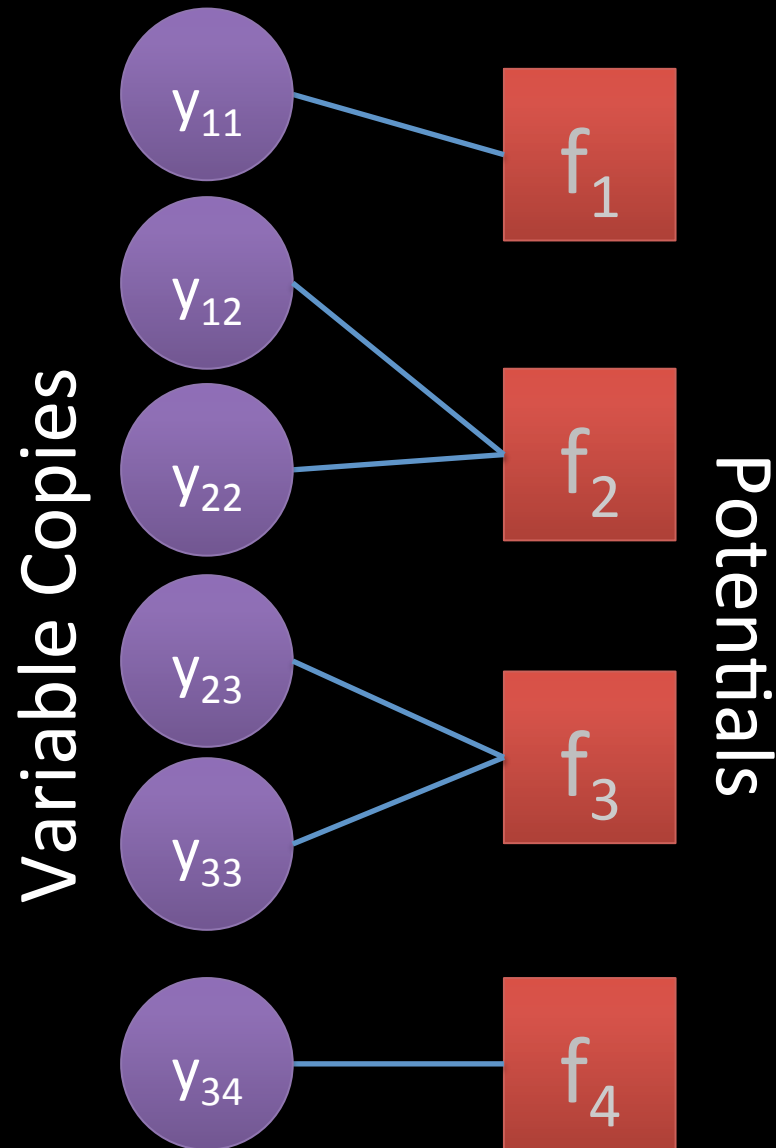
- Knapsack: combinatorial, regrets/costs, budget
- Theory: fix variables that won't change
- Practice: how can we know what will change?
- Idea: Can we use features of past inferences?
- Explore optimization (case study ADMM & PSL)

ADMM Inference in PSL

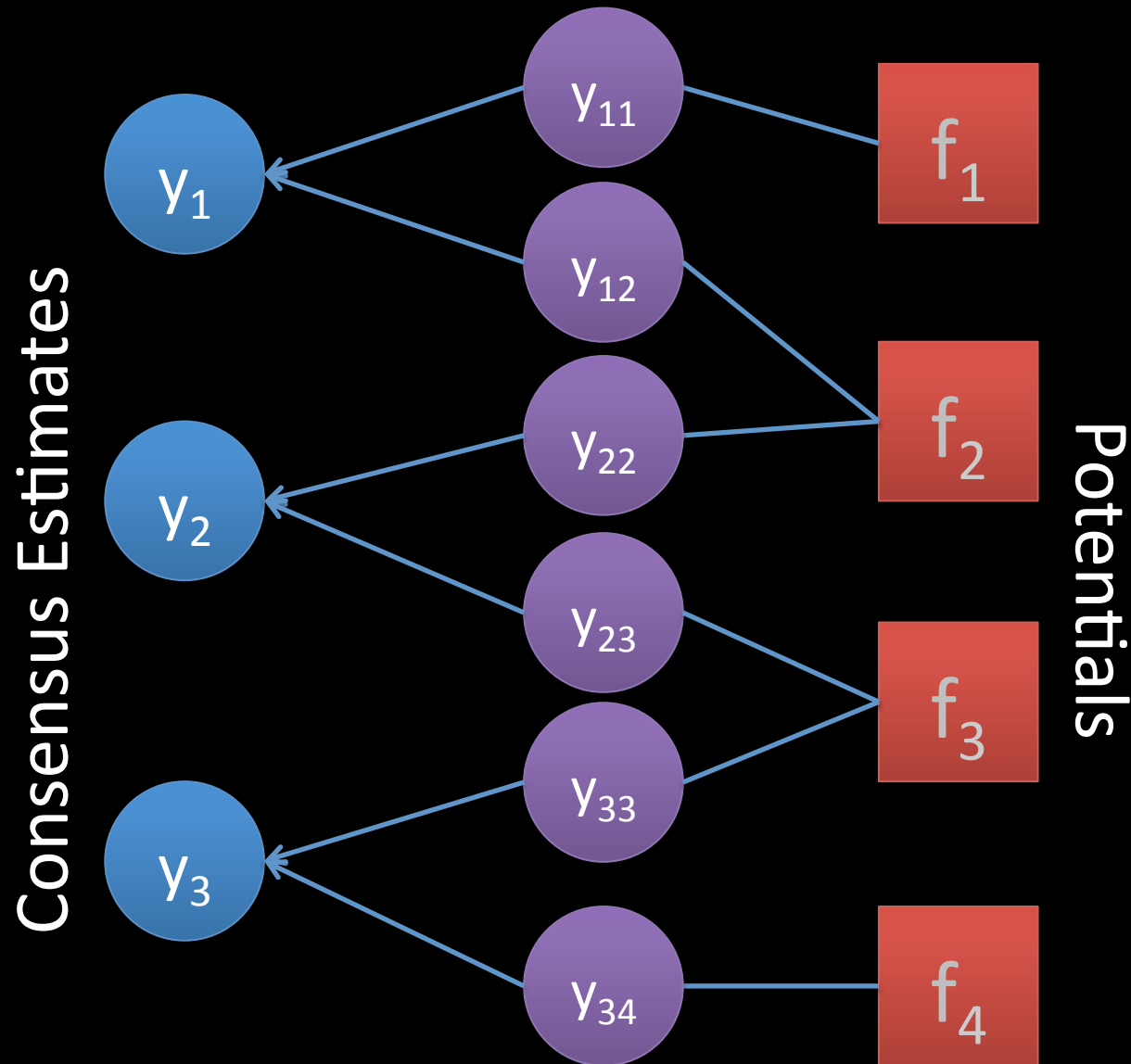
(Boyd et al., 2011; Bach et al. 2012)



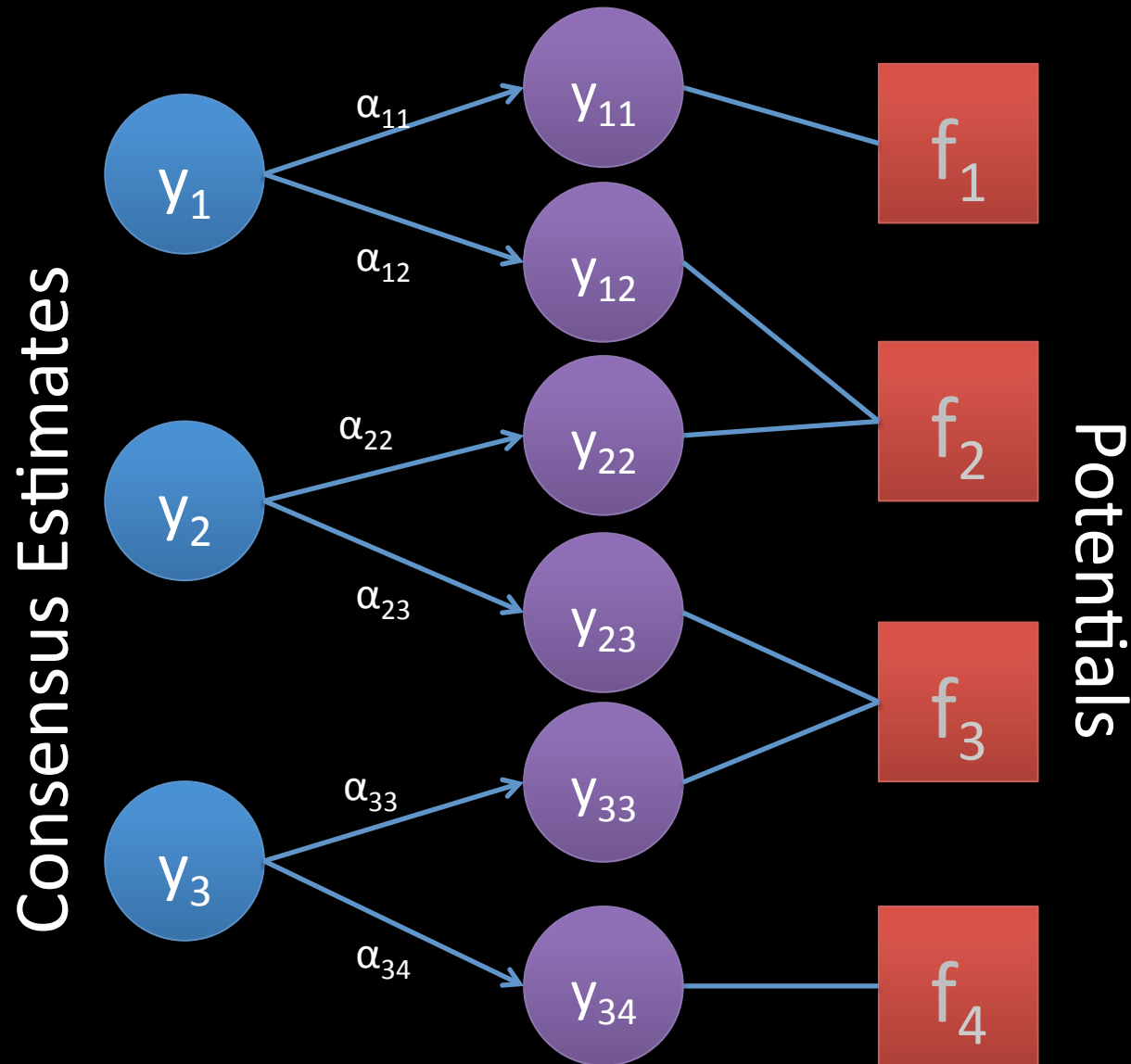
ADMM Inference in PSL



ADMM Inference in PSL



ADMM Inference in PSL



ADMM Features

$$\min_{\tilde{\mathbf{y}}_g} w_g f_g(\mathbf{x}, \tilde{\mathbf{y}}_g) + \frac{\rho}{2} \left\| \tilde{\mathbf{y}}_g - \mathbf{y}_g + \frac{1}{\rho} \boldsymbol{\alpha}_g \right\|_2^2$$

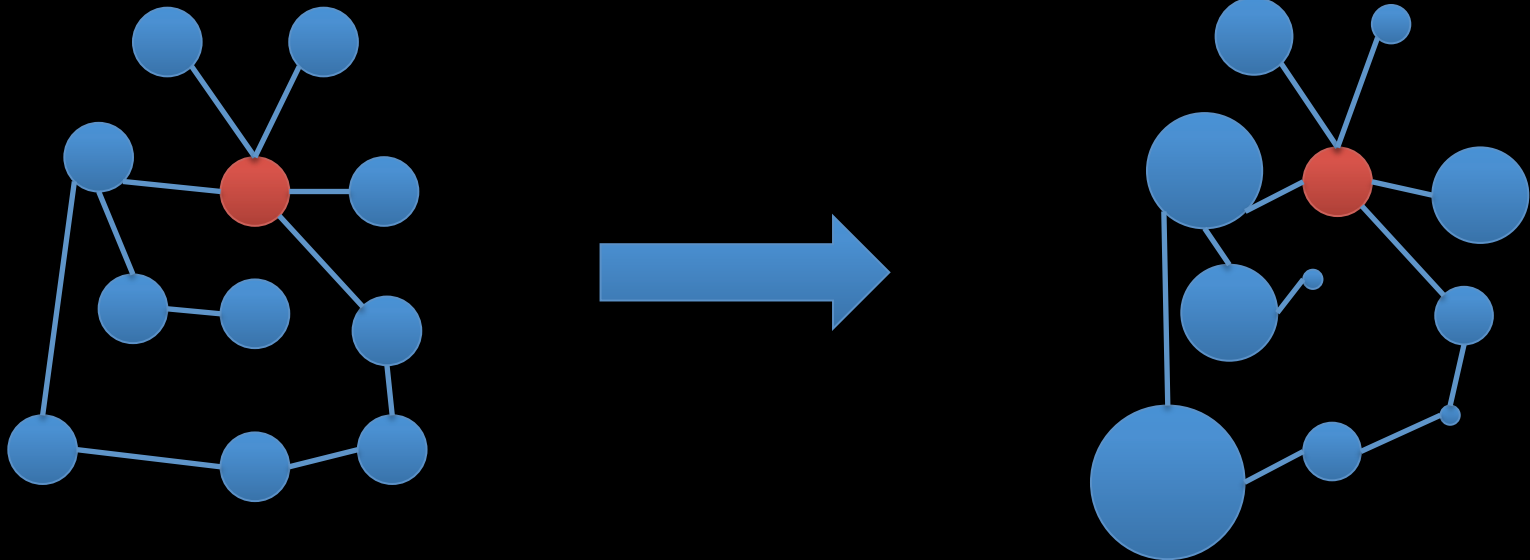
- Weight: how important is the potential?
- Potential: what loss do we incur?
- Consensus: what is the variable's value?
- Lagrange Multiplier: how much disagreement is there across potentials?

Two heuristics for activation

- **Truth-Value:** Variable value near 0.5
- **Weighted Lagrangian:** rule weight x Lagrange multipliers high

Using Model Structure

- Variable dependencies matter!
- Perform BFS, starting with new evidence
- Use heuristics + decay to prioritize exploration



EXPERIMENTAL EVALUATION

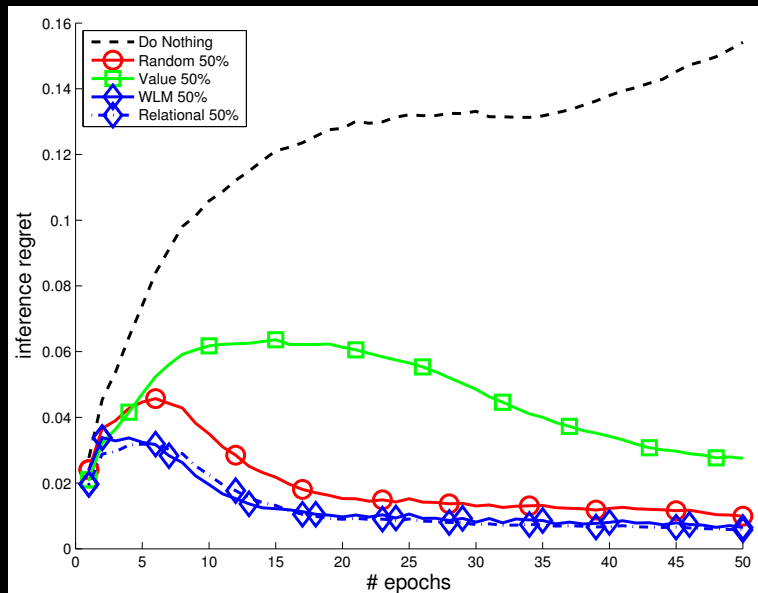
Two Online Inference Tasks

- Collective Classification (Synthetic)
 - Infer attributes of users in a social network as progressively more information is shared
- Collaborative Filtering (Jester; Goldberg et al. 2001)
 - Infer user ratings of jokes as users provide ratings for an increasing number of jokes

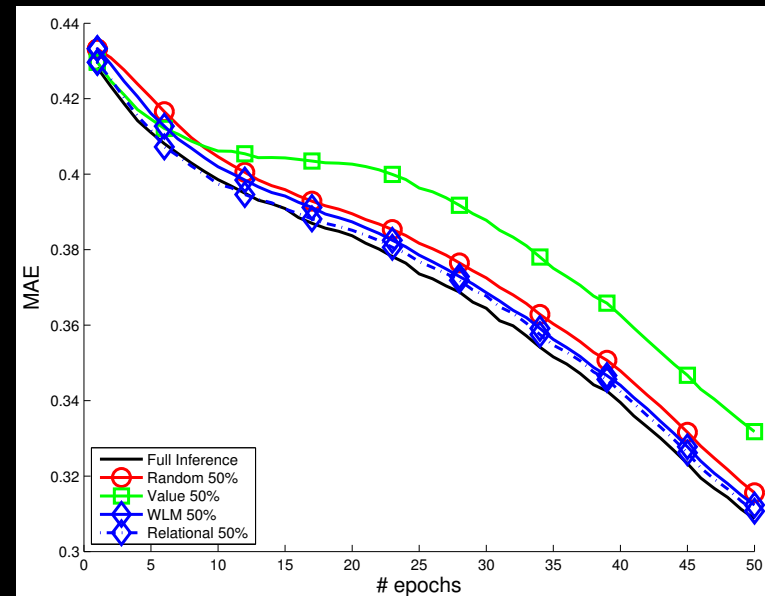
Two Online Inference Tasks

- Collective Classification (Synthetic)
 - 100 total trials (10 networks x 10 series)
 - Network evolves from 10% to 60% observed
 - Fix 50% of variables at each epoch
- Collaborative Filtering (Jester)
 - 10 trials, 100 users, 100 jokes
 - Evolve from 25% to 75% revealed ratings
 - Fix {25,50,75}% of variables at each epoch

Collective Classification: Approximate Inference



regret vs. epochs

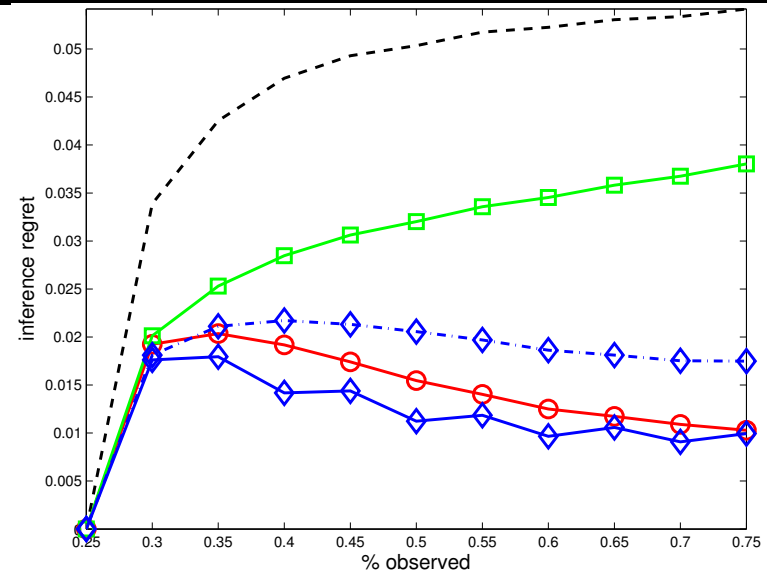
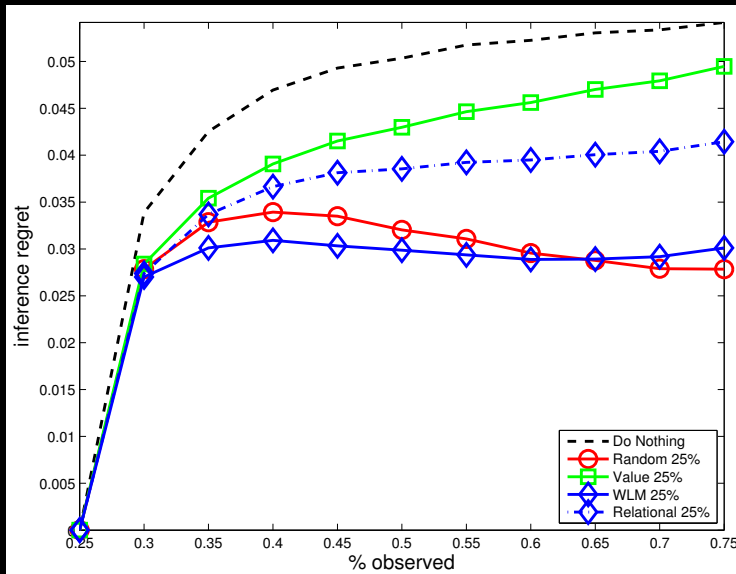


error vs. epochs

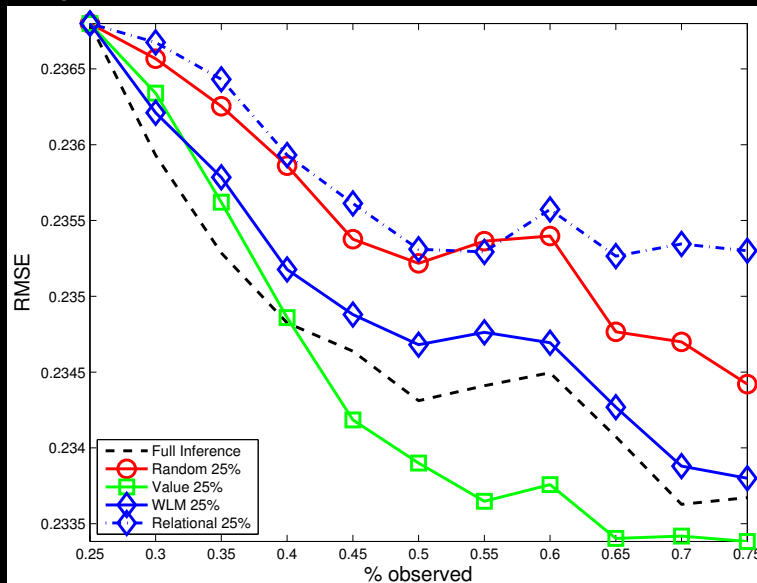
- Regret diminishes over time
- Error decreases, approaching full inference
- 69% reduction in inference time

Collaborative Filtering

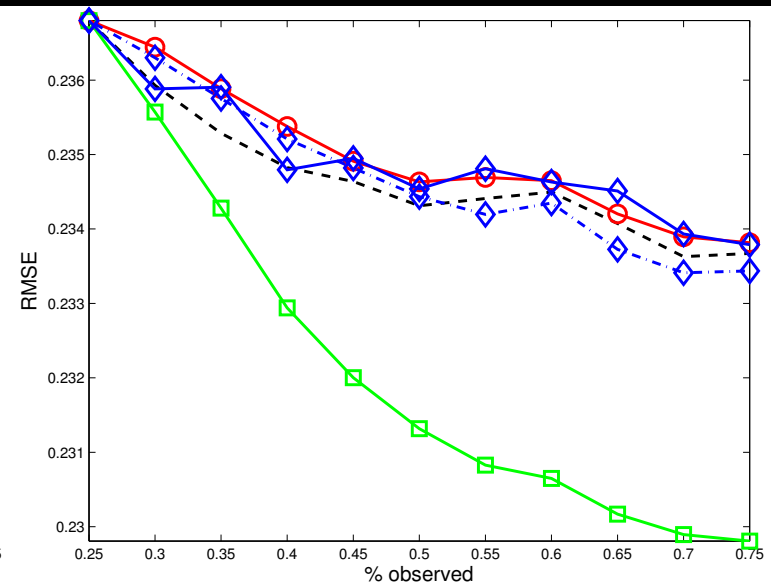
Regret



Epochs 25% activated



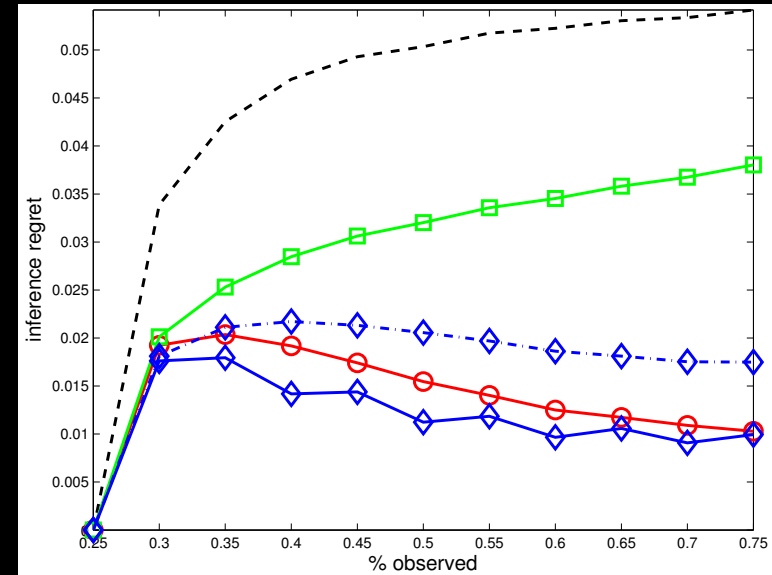
50% activated



RMSE

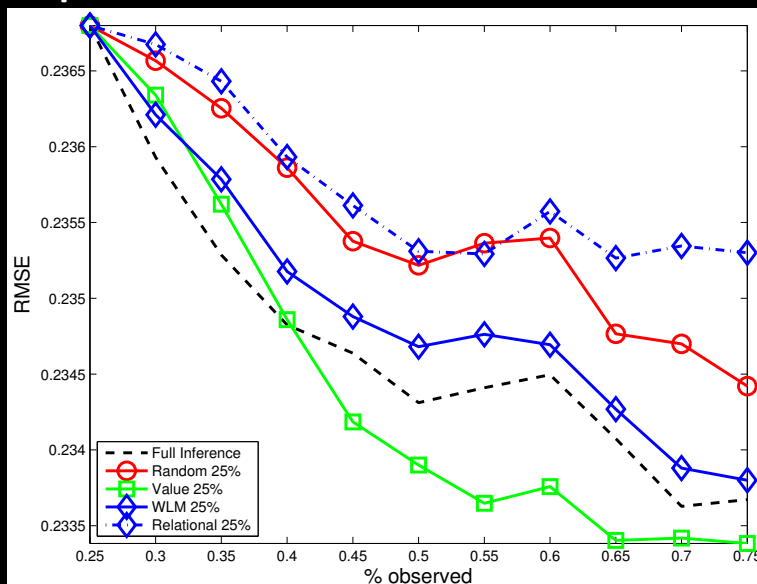
Collaborative Filtering

- **Value**: high regret, but lower error than full inference
 - Preserves polarized ratings
- 66% reduction in time for approximate inference

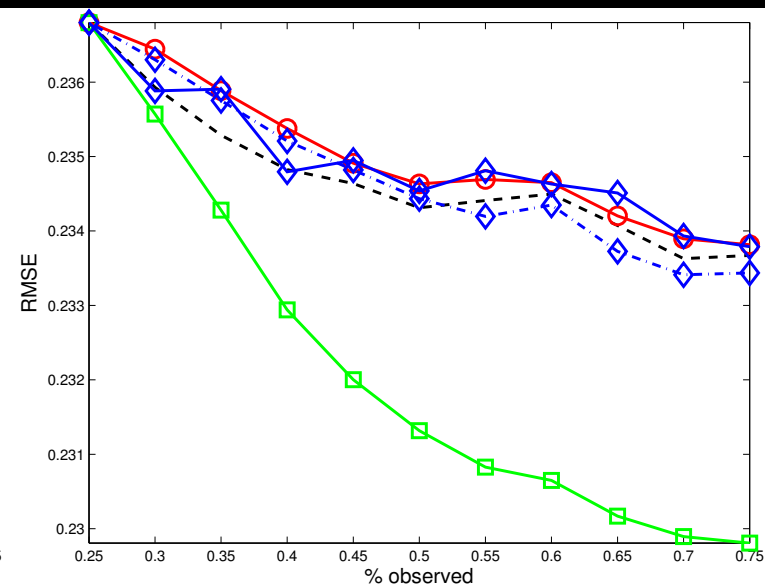


RMSE

Epochs 25% activated



50% activated



Online Collective Inference

CONCLUSION

Summary

- Extremely relevant to modern problems
- Necessity: approximate MAP state in PGMs
- Inference regret: bound approximation error
- Approx. algos: use optimization features
- Results: low regret, low error, faster
- New possibilities: rich models, fast inference

Future Work

- Better bounds for approximate inference?
- Dealing with changing models/weights
- Explicitly modeling change in models
- Applications:
 - Drug targeting
 - Knowledge Graph construction
 - Context-aware mobile devices