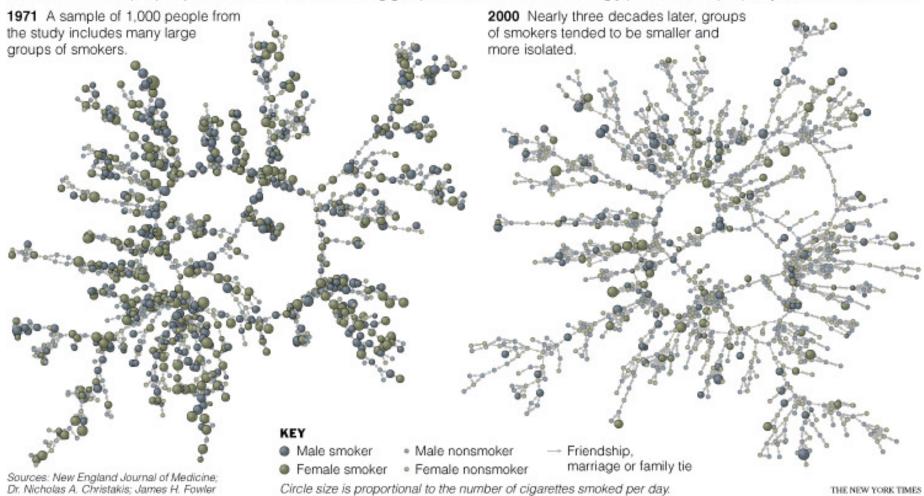
# Fully Parallel Inference in Markov Logic Networks

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#### Smoking and Quitting in Groups

Researchers studying a network of 12,067 people found that smokers and nonsmokers tended to cluster in groups of close friends and family members. As more people quit over the decades, remaining groups of smokers were increasingly pushed to the periphery of the social network.



Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

# Smokes

Name	Value
Anna	yes

# Cancer

Name	Value
Anna	no

Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

### **Smokes**

Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1: 1.5 \ \forall x. Smokes(x) \Rightarrow Cancer(x)$ 

Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

## **Smokes**

Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1: 1.5 \ \forall x. Smokes(x) \Rightarrow Cancer(x)$ 

 $F_2$ : 1.1  $\forall x. \forall y. Friends(x, y) \Rightarrow (Smokes(x) \Leftrightarrow Smokes(y))$ 

F(A,B)

F(A,A)

S(A)

S(B)

F(B,B)

C(A)

F(B,A)

C(B)

Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

# **Smokes**

Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1: 1.5 \ \forall x. Smokes(x) \Rightarrow Cancer(x)$ 

 $F_2$ : 1.1  $\forall x. \forall y. Friends(x, y) \Rightarrow (Smokes(x) \Leftrightarrow Smokes(y))$ 

True

False

Unknown

F(A,B)

F(A,A)

S(A)

S(B)

F(B,B)

C(A)

F(B,A)

C(B)

Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

## **Smokes**

Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1: 1.5 \ \forall x. Smokes(x) \Rightarrow Cancer(x)$ 



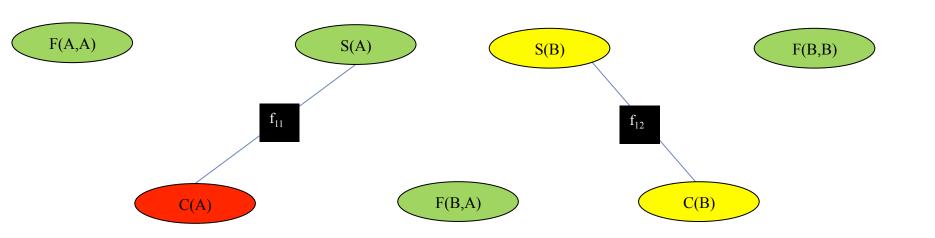
 $F_2$ : 1.1  $\forall x. \forall y. Friends(x, y) \Rightarrow (Smokes(x) \Leftrightarrow Smokes(y))$ 

True

False

Unknown





Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

### **Smokes**

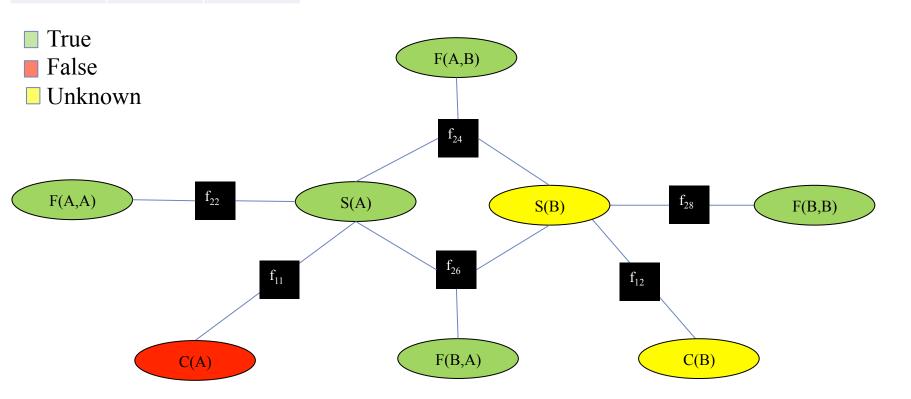
Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1: 1.5 \ \forall x. Smokes(x) \Rightarrow Cancer(x)$ 





Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

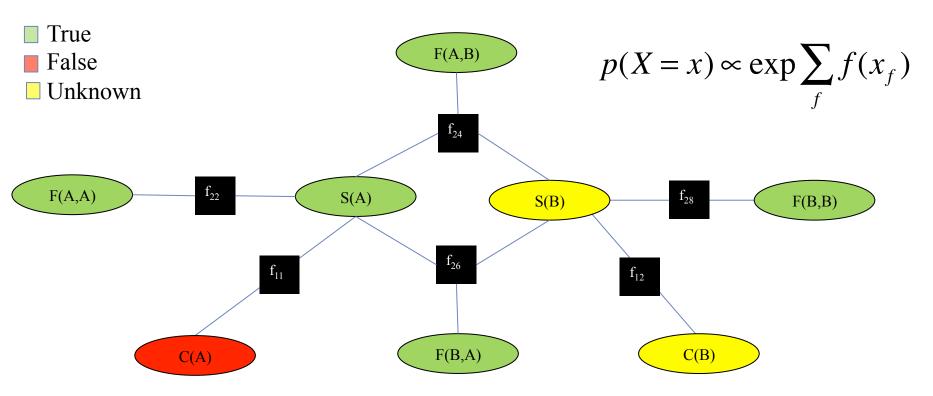
### **Smokes**

Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1$ : 1.5  $\forall x$ .Smokes $(x) \Rightarrow$  Cancer(x)



Name1	Name2	Value
Anna	Bob	yes
Bob	Anna	yes
Anna	Anna	yes
Bob	Bob	yes

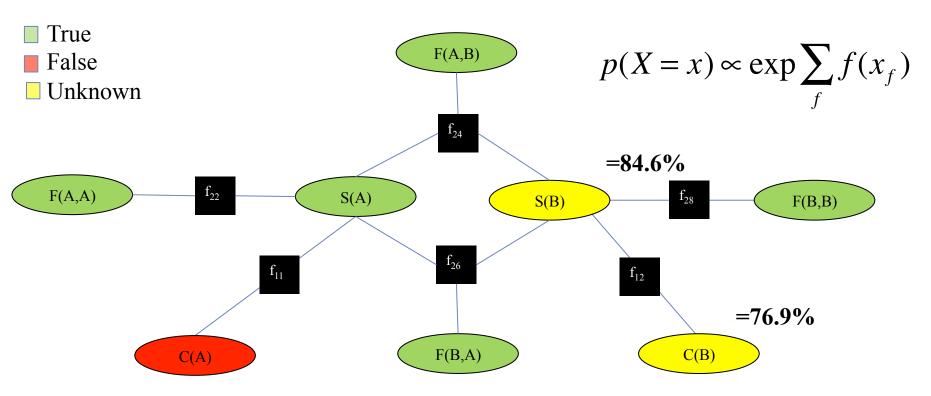
### **Smokes**

Name	Value
Anna	yes

### Cancer

Name	Value
Anna	no

 $F_1$ : 1.5  $\forall x$ .Smokes $(x) \Rightarrow Cancer(x)$ 



# Inference in Markov Logic Networks (I)

### Sampling in MNL

- Approximation unavoidable
- Generic technique to approximate expectations
- Simple, versatile, well understood

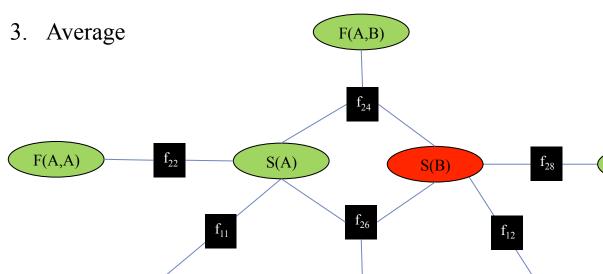
# Inference in Markov Logic Networks (I)

#### Sampling in MNL

- Approximation unavoidable
- Generic technique to approximate expectations
- Simple, versatile, well understood

#### Sampling process

- 1. Assign a value to each variable
- 2. Count



F(B,A)

Var	#true	#false
F(A,A)	XX	XX
F(A,B)	XX	XX
F(B,A)	XX	XX
F(B,B)	XX	XX
S(A)	XX	XX
S(B)	XX	XX
C(A)	XX	XX
C(B)	XX	XX

F(B,B)

# Inference in Markov Logic Networks (I)

#### Sampling in MNL

- Approximation unavoidable
- Generic technique to approximate expectations
- Simple, versatile, well understood

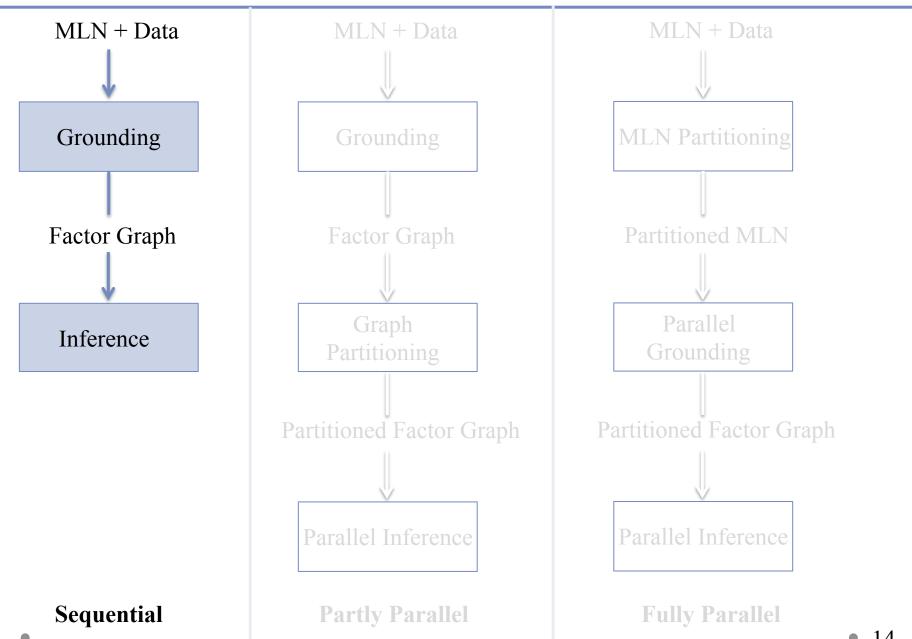
#### Sampling process

- 1. Assign a value to each variable
- 2. Count
- 3. Average

$$\frac{1}{n} \sum_{i=1}^{n} h(x^{(i)}) = \hat{\mu}$$
  $Var_p[\hat{\mu}] = \frac{Var_p[h(x)]}{n}$ 

More samples more efficiency

# Sequential approach



# Networks can be very large

### Lots of applications

- Link prediction
- Information Extraction
- Entity Resolution
- Ontology Learning

#### How to gain scalability?

- Grounding is expensive
- Inference is expensive

### Networks can be very large

#### Lots of applications

- Link prediction
- Information Extraction
- Entity Resolution
- Ontology Learning

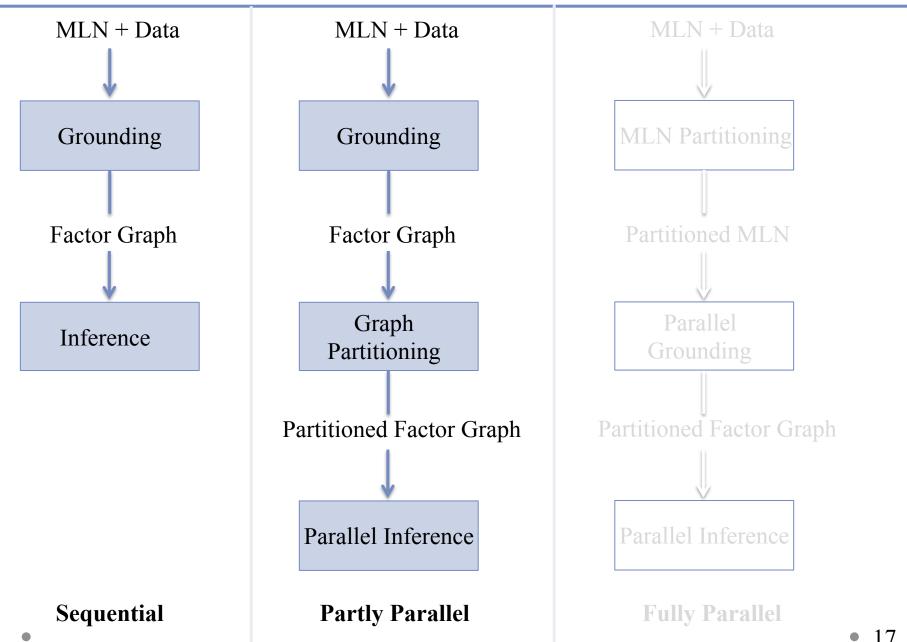
#### How to gain scalability?

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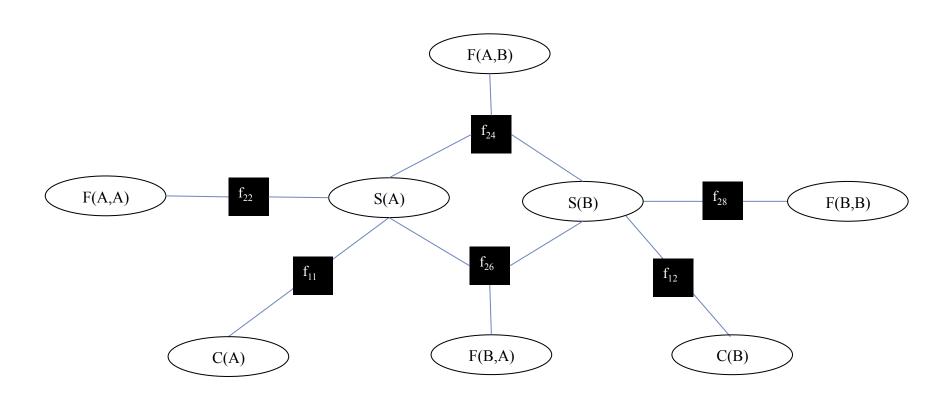
### Why speed up sampling?

- Expensive
- Datasets can be big
- Dataset 72k variables each sample between 2-5 seconds
- 1 million samples  $\approx 50$  days

# Partly parallel approach

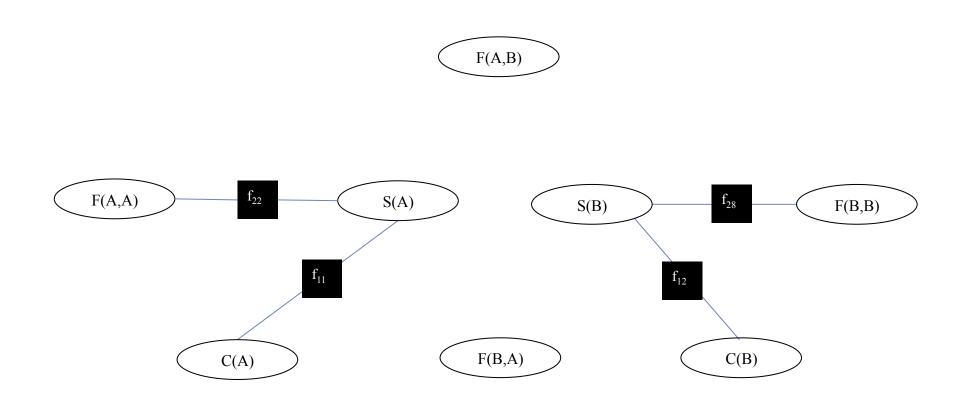


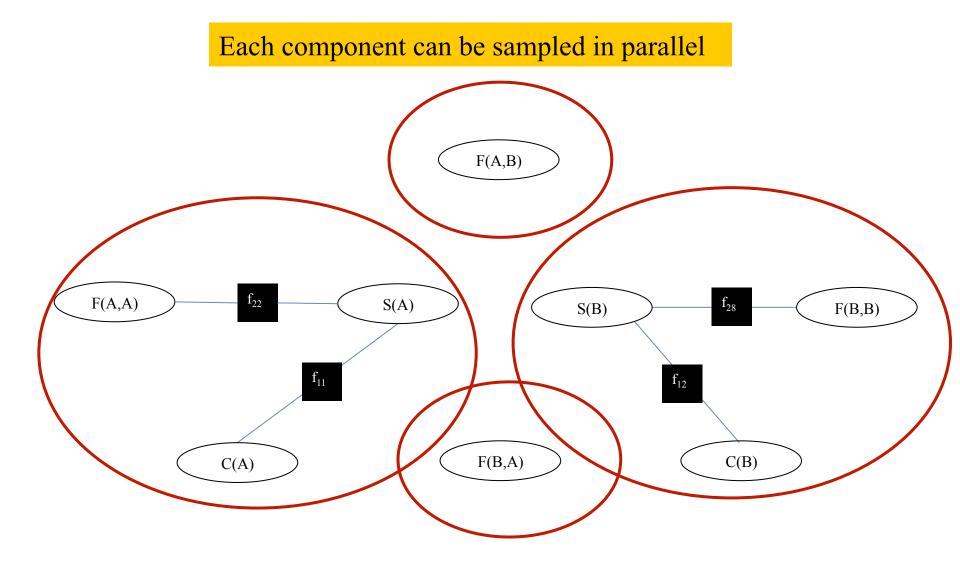
Cut the network to sample each partition in parallel

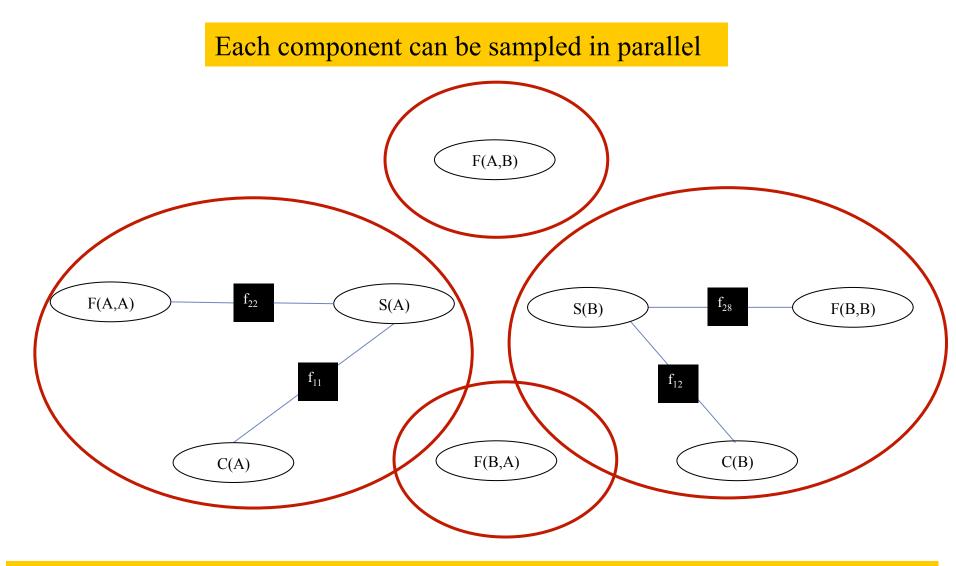


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Cut is performed by removing factors to generate independent components



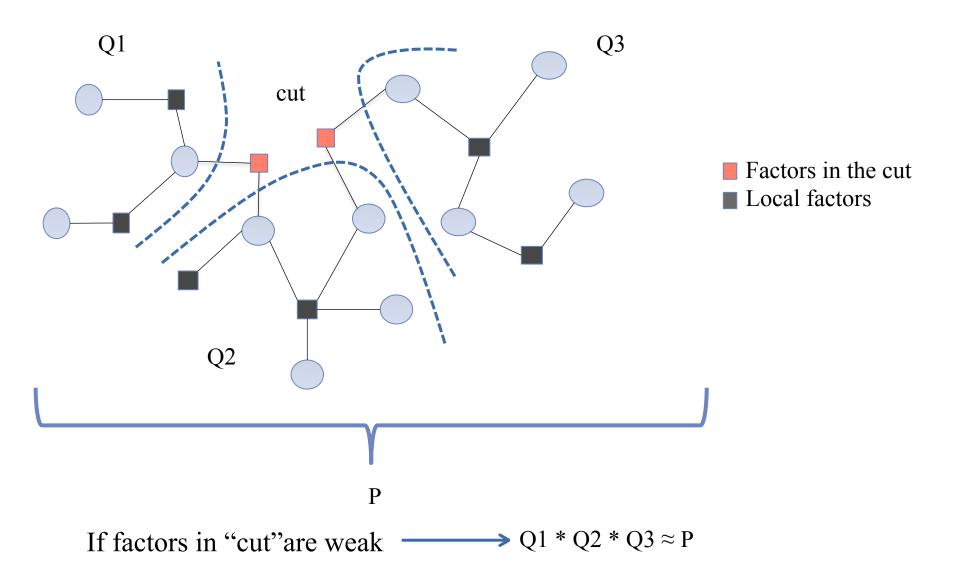


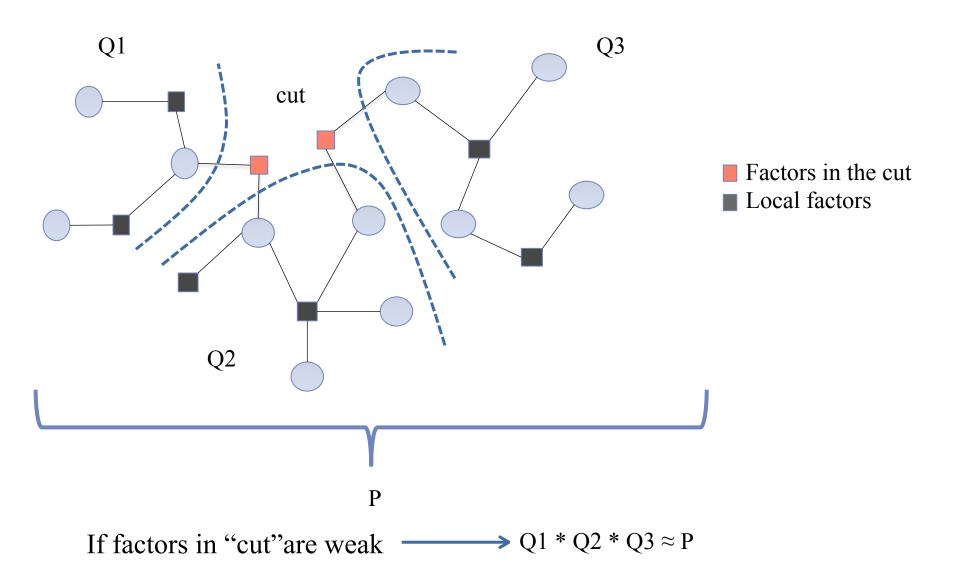


Information loss equivalent to lost connections. How big is the information loss?

### Outline

- Background and Motivation
- Parallel Inference
- Parallel Grounding
- Conclusion



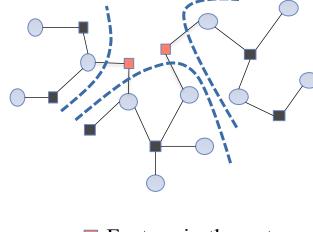


How to find a cut with weak factors?

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# Importance Sampling

- 1. Cut the graph to get independent components
- 2. Get a sample from each component independently
- 3. Correct the sample to match the original distribution
- 4. Correction determined by factors in cut



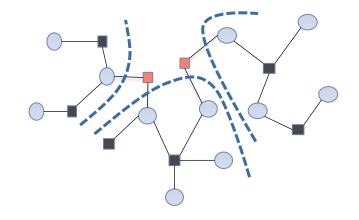
Factors in the cut

Local factors

Efficiency of the estimation depends on the information loss (factors in the cut)

# Importance Sampling

- 1. Cut the graph to get independent components
- 2. Get a sample from each component independently
- 3. Correct the sample to match the original distribution
- 4. Correction determined by factors in cut



- Factors in the cut
- Local factors

Efficiency of the estimation depends on the information loss (factors in the cut)

### Standard Monte-Carlo

$$Var_p[\hat{\mu}] = \frac{Var_p[h(x)]}{n}$$

# Importance Sampling

$$Var_q(\hat{\mu}_{is}) \approx \frac{(1 + Var_q[w(x)])Var_p[h(x)]}{n}$$

w(x): sum of the instantiated factors in the cut for a sample

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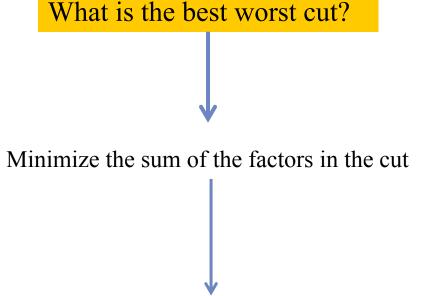
- (

# Quality of the cut: a bound

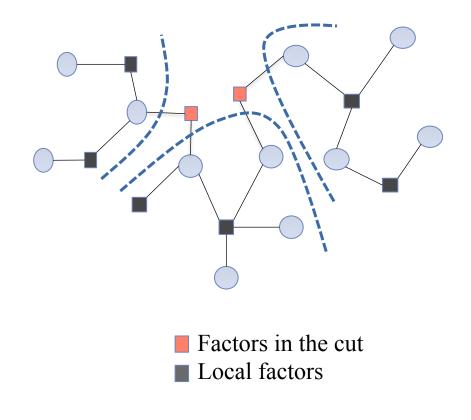
- Calculating the dispersion of the weights in the cut is intractable
- What is the worst possible quality for each cut?
- What is the best of the worst?

# Quality of the cut: a bound

- Calculating the dispersion of the weights in the cut is intractable
- What is the worst possible quality for each cut?

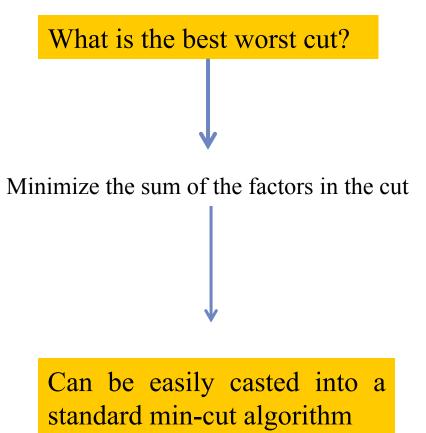


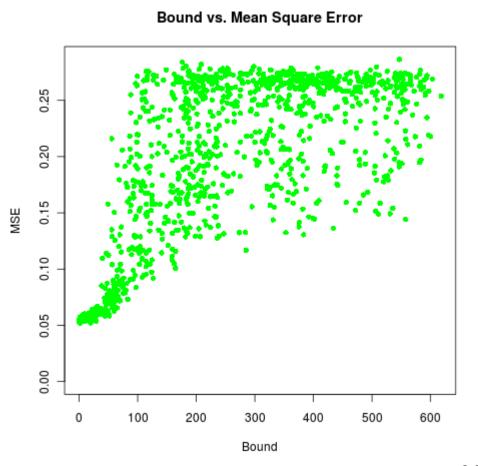
Can be easily casted into a standard min-cut algorithm



# Quality of the cut: a bound

- Calculating the dispersion of the weights in the cut is intractable
- What is the worst possible quality for each cut?





# Results Parallel Inference with Importance Sampling

#### Dataset

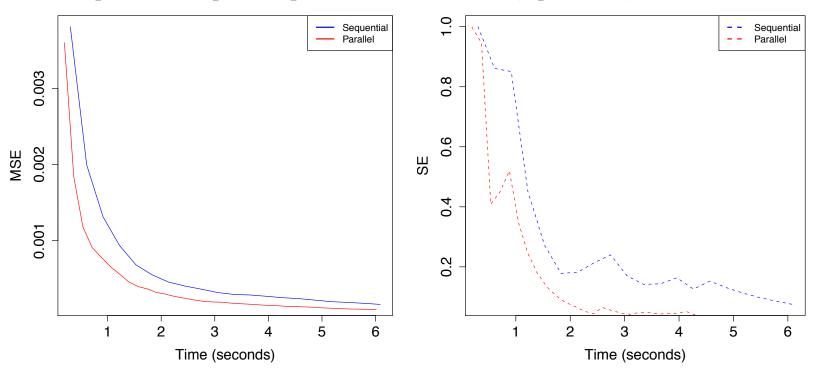
- UW-CSE (22 predicates, 94 clauses)
  - Link prediction
- ~9K variables and ~1M factors (after grounding)

# Results Parallel Inference with Importance Sampling

#### Dataset

- UW-CSE (22 predicates, 94 clauses)
  - Link prediction
- ~9K variables and ~1M factors (after grounding)

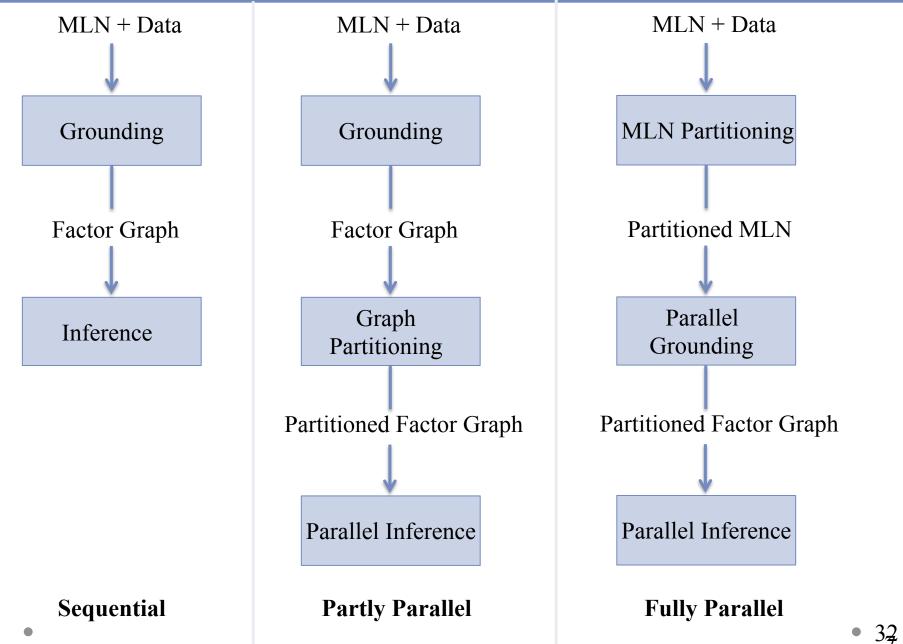
Sequential and parallel probabilistic inference (4 partitions)



**Average MSE** 

**Maximum SE** 

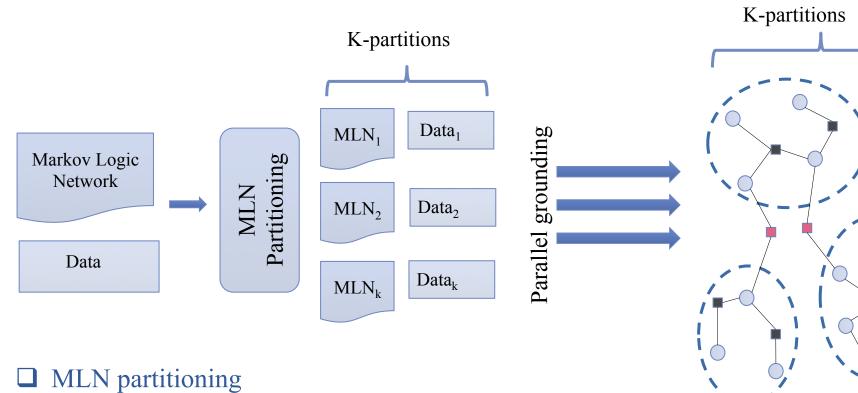
# Fully Parallel Approach



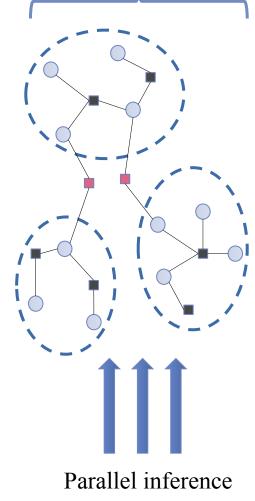
### Outline

- Background and Motivation
- Parallel Inference
- Parallel Grounding
- Conclusion

# Parallel Grounding



- Use information at the schema level
- Compute partitions before grounding
- Ground partitions in parallel
- Avoids expensive graph cuts



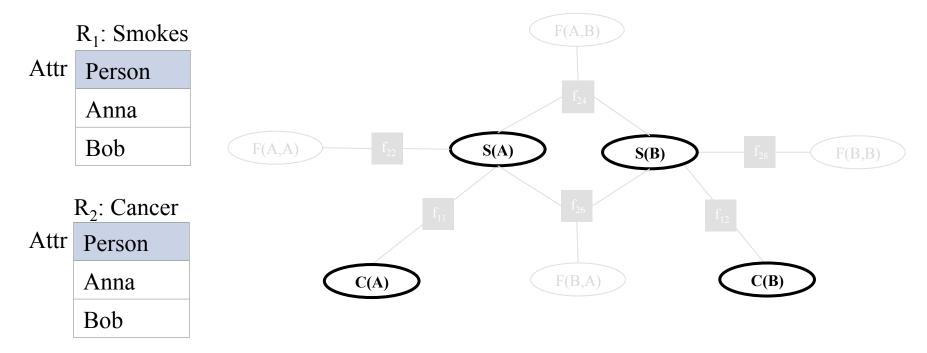
# Grounding $\equiv$ Database joins

Formula $Smokes(x) \Rightarrow Cancer(x)$	$CNF \neg Smokes(x) \lor Cancer(x)$
Predicates and domain	Ground Clauses
Smokes(person)	$\neg Smokes(Anna) \lor Cancer(Anna)$
Cancer( $person$ ) $person = \{Anna, Bob\}$	$\neg Smokes(Bob) \lor Cancer(Bob)$

# Grounding $\equiv$ Database joins

Formula $Smokes(x) \Rightarrow Cancer(x)$	$CNF \neg Smokes(x) \lor Cancer(x)$
Predicates and domain	Ground Clauses
Smokes(person)	$\neg Smokes(Anna) \lor Cancer(Anna)$
Cancer( $person$ ) $person = \{Anna, Bob\}$	$\neg Smokes(Bob) \lor Cancer(Bob)$

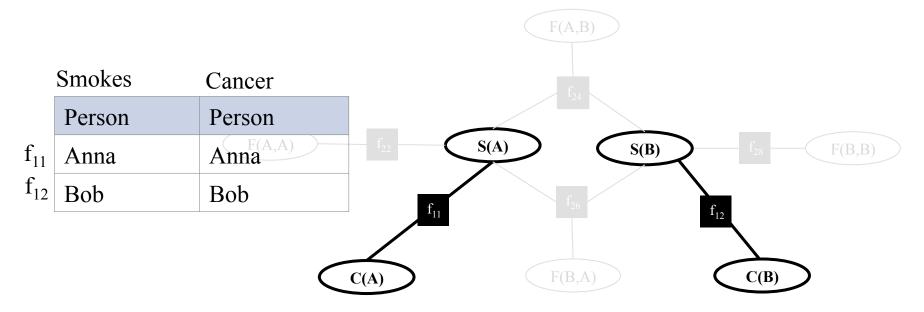
Ground variables corresponds to Relations



# Grounding $\equiv$ Database joins

Formula $Smokes(x) \Rightarrow Cancer(x)$	$CNF \neg Smokes(x) \lor Cancer(x)$
Predicates and domain	Ground Clauses
Smokes( $person$ ) Cancer( $person$ ) $person = \{Anna, Bob\}$	$\mathbf{f_{11}} \neg \operatorname{Smokes}(Anna) \vee \operatorname{Cancer}(Anna)$ $\mathbf{f_{12}} \neg \operatorname{Smokes}(Bob) \vee \operatorname{Cancer}(Bob)$

• Ground clauses corresponds to natural join: Smokes ⋈ Cancer



```
\mathbf{R}(x,y)

\mathrm{Dom}(x) = \{\mathrm{Anna, Bob}\}

\mathrm{Dom}(y) = \{\mathrm{Charles, Debbie}\}
```

### Attributes

K	
X	y
Anna	Charles
Bob	Charles
Anna	Debbie
Bob	Debbie

R(A,C)



R(B,D)

 $\mathbf{R}(x,y)$   $\mathrm{Dom}(x) = \{\mathrm{Anna, Bob}\}$  $\mathrm{Dom}(y) = \{\mathrm{Charles, Debbie}\}$ 

$$R_1 = \sigma_{y=Charles}(R)$$

 $R_2 = \sigma_{y=\underline{Debbie}}(R)$ 

Attributes

$R_1$		
X	y	
Anna	Charles	
Bob	Charles	

Attributes

 $R_2$ 

X	y
Anna	Debbie
Bob	Debbie

R(A, C)



$$(R(A, \mathbf{D}))$$

 $R(B, \mathbf{D})$ 

```
\mathbf{R}(x,y)

\mathrm{Dom}(x) = \{\mathrm{Anna, Bob}\}

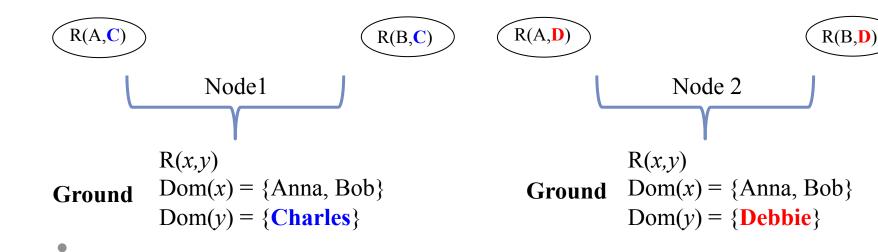
\mathrm{Dom}(y) = \{\mathrm{Charles, Debbie}\}
```

$$R_1 = \sigma_{v=Charles}(R)$$

 $R_2 = \sigma_{y=\underline{Debbie}}(R)$ 

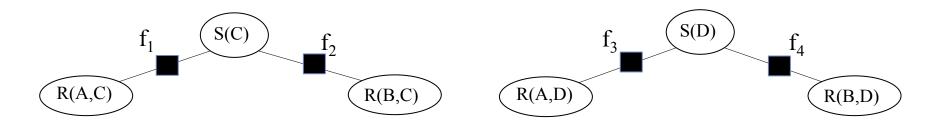
	$R_1$	
Attributes	X	У
	Anna	Charles
	Bob	Charles

	$R_2$	
Attributes	X	У
	Anna	Debbie
	Bob	Debbie



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**f**:  $R(x,y) \lor S(y)$   $Dom(x) = \{Anna, Bob\}$  $Dom(y) = \{Charles, Debbie\}$ 



f:  $R(x,y) \lor S(y)$   $Dom(x) = \{Anna, Bob\}$  $Dom(y) = \{Charles, Debbie\}$ 

	R		S
	X	y	y
$\mathbf{f}_1$	Anna	Charles	Charles
$\mathbf{f_2}$	Bob	Charles	Charles
$\mathbf{f_3}$	Anna	Debbie	Debbie
$\mathbf{f_4}$	Bob	Debbie	Debbie
		$R \bowtie_{R,y=S,y}$	S



f: 
$$R(x,y) \lor S(y)$$
  
 $Dom(x) = \{Anna, Bob\}$   
 $Dom(y) = \{Charles, Debbie\}$ 

$$R_{1} = \sigma_{y=Charles}(R)$$

$$S_{1} = \sigma_{y=Charles}(S)$$

 $R_1$ 

 $S_1$ 

	X	У	у
<b>f</b> <sub>1</sub>	Anna	Charles	Charles
$\mathbf{f_2}$	Bob	Charles	Charles

 $R_1 \bowtie S_1 \atop R_1.y = S_1.y$ 

$R_2$	=	$\sigma_{v=Debbie}$	(R)
-------	---	---------------------	-----

$$S_2 = \sigma_{y=Debbie}(S)$$

 $R_2$ 

 $S_2$ 

	X	у	y
$\mathbf{f}_3$	Anna	Debbie	Debbie
$\mathbf{f_4}$	Bob	Debbie	Debbie

$$R_2 \underset{\mathbf{R_2.y}=\mathbf{S_2.y}}{\bowtie} S_2$$

Local join

$$\mathbf{f}: \mathbf{R}(x,y) \vee \mathbf{S}(y)$$

 $Dom(x) = \{Anna, Bob\}$ 

 $Dom(y) = \{Charles, Debbie\}$ 

$$R_1 = \sigma_{y=Charles}(R)$$

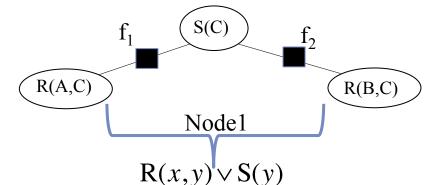
$$S_1 = \sigma_{y=Charles}(S)$$

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		- 1	

 $\mathbf{S}_1$ 

	X	У	У
1	Anna	Charles	Charles
2	Bob	Charles	Charles

$$R_1 \bowtie S_1$$
 $R_1 v = S_1 v$ 



Ground

 $Dom(x) = \{Anna, Bob\}$ 

 $Dom(y) = \{ \frac{\textbf{Charles}}{\textbf{Charles}} \}$ 

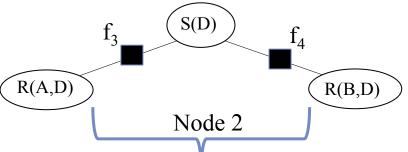
$$R_2 = \sigma_{y=\underline{Debbie}}(R)$$

$$S_2 = \sigma_{v=Debbie}(S)$$

R<sub>2</sub>

	X	У	У	
$\mathbf{f}_3$	Anna	Debbie	Debbie	
$\mathbf{f}_4$	Bob	Debbie	Debbie	





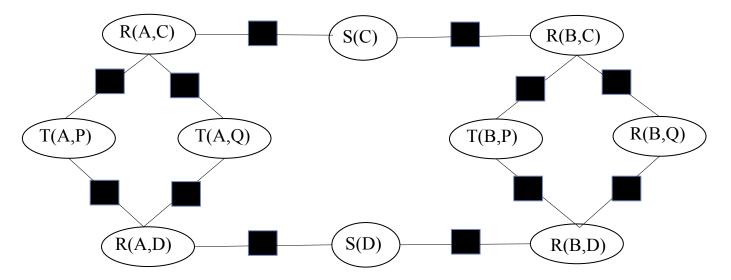
 $R(x,y) \vee S(y)$ 

**Ground**  $Dom(x) = \{Anna, Bob\}$ 

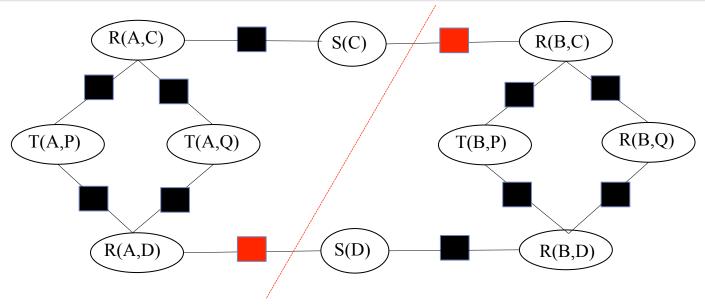
 $Dom(y) = \{ \mathbf{Debbie} \}$ 

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```
f_1: R(x,y) \vee S(y)
f_2: R(x,y) \vee T(x,z)
Dom(x) = \{A, B\}
Dom(y) = \{C, D\}
Dom(z) = \{P, Q\}
```



```
f_1: R(x,y) \vee S(y)
f_2: R(x,y) \vee T(x,z)
Dom(x) = \{A, B\}
Dom(y) = \{C, D\}
Dom(z) = \{P, Q\}
```



How to compute partitions at the Markov logic level?

$$f_1: R(x,y) \vee S(y)$$

$$f_2: R(x,y) \vee T(x,z)$$

$$Dom(x) = \{A, B\}$$

$$Dom(y) = \{C, D\}$$

$$Dom(z) = \{P, Q\}$$

$$S(y) \xrightarrow{J_1} R(x,y) \xrightarrow{J_2} T(x,z)$$

### Partitioning at Rule level

Model MLN as a join graph

```
f_1: R(x,y) \vee S(y)
f_2: R(x,y) \vee T(x,z)
Dom(x) = \{A, B\}
Dom(y) = \{C, D\}
Dom(z) = \{P, Q\}
```

$$S(\mathbf{y}) \xrightarrow{J_1} R(\mathbf{x}, \mathbf{y}) \xrightarrow{J_2} T(\mathbf{x}, z)$$

Partitioning at Rule level Co-partitioning strategy?

Model MLN as a join graph

```
f_1: R(x,y) \vee S(y)
f_2: R(x,y) \vee T(x,z)
Dom(x) = \{A, B\}
Dom(y) = \{C, D\}
Dom(z) = \{P, Q\}
```

$$S(y) \xrightarrow{J_1} \mathbb{R}_{xy=S,y} \mathbb{R}(x,y) \xrightarrow{J_2} \mathbb{R}_{xz=T,x} \mathbb{T}(x,z)$$

$$|J_1| = 4 \qquad |J_2| = 8$$

### Partitioning at Rule level

- Model MLN as a join graph
- Estimate join sizes

```
f_1: R(x,y) \vee S(y)
f_2: R(x,y) \vee T(x,z)
Dom(x) = \{A, B\}
Dom(y) = \{C, D\}
Dom(z) = \{P, Q\}
```

$$S(y) \xrightarrow{J_1} \mathbb{R}_{xy=S,y} \mathbb{R}(x,y) \xrightarrow{J_2} \mathbb{R}_{xz=T,x} \mathbb{T}(x,z)$$

$$|J_1| = 4 \qquad |J_2| = 8$$

### Partitioning at Rule level

- Model MLN as a join graph
- Estimate join sizes
- Co-partition to maximize size of local joins optimization problem
- Encode as an ILP

$$f_1 : R(x,y) \lor S(y)$$
  $Dom(x) = \{A, B\}$   
 $f_2 : R(x,y) \lor T(x,z)$   $Dom(y) = \{C, D\}$   
 $Dom(z) = \{P, Q\}$ 

$$S(y) \xrightarrow{J_1} R(x,y) \xrightarrow{J_2} T(x,z)$$

- $JS(J_1) = 4$
- $JS(J_2) = 8$
- Co-partition R and T on Dom(*x*)
- S on Dom(y)

### Local join

#### Ground

$$R(x,y) Dom(x) = \{A\}, Dom(y) = \{C,D\}$$
  
 $T(x,z) Dom(x) = \{A\}, Dom(z) = \{P,Q\}$   
 $S(y) Dom(y) = \{C\}$ 

#### Ground

$$R(x,y) Dom(x) = \{B\}, Dom(y) = \{C,D\}$$
  
 $T(x,z) Dom(x) = \{B\}, Dom(z) = \{P,Q\}$   
 $S(y) Dom(y) = \{D\}$ 

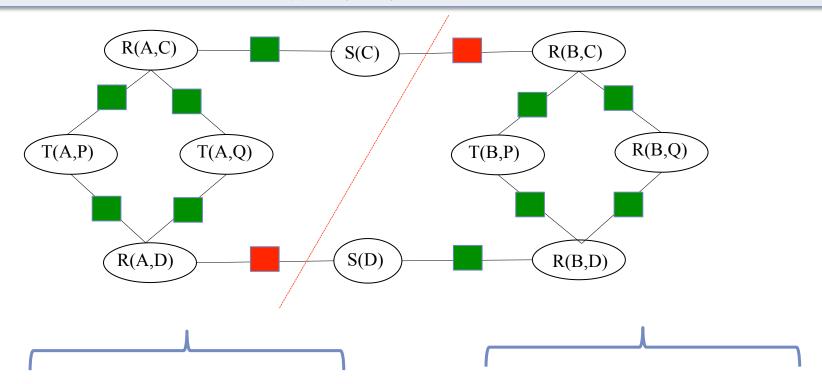
$$f_1: R(x,y) \vee S(y)$$

$$f_2: R(x,y) \vee T(x,z)$$

$$Dom(x) = \{A, B\}$$

$$Dom(y) = \{C, D\}$$

$$Dom(z) = \{P, Q\}$$



#### Ground

$$R(x,y) Dom(x) = \{A\}, Dom(y) = \{C,D\}$$
  
 $T(x,z) Dom(x) = \{A\}, Dom(z) = \{P,Q\}$   
 $S(y) Dom(y) = \{C\}$ 

#### Ground

$$R(x,y) Dom(x) = \{B\}, Dom(y) = \{C,D\}$$
  
 $T(x,z) Dom(x) = \{B\}, Dom(z) = \{P,Q\}$   
 $S(y) Dom(y) = \{D\}$ 

# MLN Partitioning (evaluation)

Comparison of various graph partitioning approaches for k partitions

$\overline{k}$	Approach	Factors in cut	Weight of cut	Balancing	Runtime
<i>k</i> = 2	РаТоН	4678	1109.04	0.000	948.288s
	Tuffy	4686	1108.66	0.000	1.092s
	MLN part.	4690	1109.47	0.000	0.003s
<i>k</i> = 4	РаТоН	63001	64500.40	0.012	952.254s
	Tuffy	7040	1662.46	0.000	1.288s
	MLN part.	7023	1662.84	0.000	0.003s

### Outline

- Background and Motivation
- Parallel Inference
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### Conclusions

### Markov logic networks

- Incomplete database + first order rules
- Scalability challenges

### First fully parallel approach to MLN inference

- Partition the MLN before grounding
- Ground partitions in parallel
- Run parallel inference

### Preliminary experimental results

- Orders of magnitude faster partitioning at similar quality
- Parallel inference effective

### Conclusions

# Markov logic networks Questions?

- Incomplete database + first order rules
- Scalability challenges

### First fully parallel approach to MLN inference

- Partition the MLN before grounding
- Ground partitions in parhank you!
- Run parallel inference

### Preliminary experimental results

- Orders of magnitude faster partitioning at similar quality
- Parallel inference effective