### Learning with Approximate Inference

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### **Topics**

- Learning MN parameters using Gradient Ascent and Belief Propagation
  - Difficulty with exact methods
- Approximate methods
  - Approximate Inference
    - Belief Propagation
    - Sampling-based Learning
    - MAP-based Learning
- Ex: CRF for Protein Structure Prediction

### Likelihood function for a Markov Network

• Log-linear form of MN with parameters  $\theta$  is

$$P(X_1,..X_n;\theta) = \frac{1}{Z(\theta)} \exp\left\{\sum_{i=1}^k \theta_i f_i(D_i)\right\}$$

where  $\theta = \{\theta_1, ..., \theta_k\}$  are k parameters, each associated with a feature  $f_i$  defined over instances of  $D_i$ 

Where the partition function defined as:

$$Z(\theta) = \sum_{\xi} \exp\left\{\sum_{i} \theta_{i} f_{i}(\xi)\right\}$$

• We wish to find  $\theta$  that maximizes the Log-likelihood

$$\ell(\theta:D) = \sum_{i} \theta_{i} \left( \sum_{m} f_{i} \left( \xi[m] \right) \right) - M \ln Z(\theta)$$

 $\partial \ell(\theta)$ 

### Gradient-based learning of Undirected PGM

Initialize  $\theta$ 

*Goal:* Determine parameters  $\theta$ of Markov Network over variables  $\chi$  given data set  $\mathcal{D}$  with k features  $f_i$ 

$$P_{\theta}(\chi) = \frac{1}{Z(\theta)} \exp\left[-\sum_{i=1}^{k} \theta_{i} f_{i}(D_{i})\right]$$

Run inference to compute  $Z(\theta)$ ,  $E_{\theta}[f_i]$ 

$$Z(\theta) = \sum_{\xi} \exp\left\{\sum_{i} \theta_{i} f_{i}(\xi)\right\}$$

$$Z(\theta) = \sum_{\xi} \exp\left\{\sum_{i} \theta_{i} f_{i}(\xi)\right\} \qquad E_{\theta}[f_{i}] = \frac{1}{Z(\theta)} \sum_{\xi} f_{i}(\xi) \exp\left\{\sum_{j} \theta_{j} f_{j}(\xi)\right\}$$

Compute gradient of likelihood  $\ell$ 

Update 
$$\theta$$
 with step size  $n$ 

step size n

Stop

$$\left| \frac{\partial}{\partial \theta_{i}} \ell \left( \theta; \mathcal{D} \right) = E_{\mathcal{D}}[f_{i}(\chi)] - E_{\theta}[f_{i}] \right| \quad \left| \nabla \ell(\theta) = \frac{d}{d\theta} \ell(\theta) = \left[ \frac{\partial \ell(\theta)}{\partial \theta_{i}} \right] \right|$$

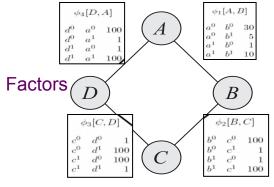
$$\boxed{\boldsymbol{\theta}^{t+1} \leftarrow \boldsymbol{\theta}^t + \eta \nabla \ell \big(\boldsymbol{\theta}^t; \boldsymbol{\mathcal{D}}\big)}$$

$$oldsymbol{ heta}^{t+1} - oldsymbol{ heta}^t \leq \delta$$

Method assumes that we are able to compute partition function  $Z(\theta)$ , which is a sum over all unnormalized probabilities of assignments  $\xi$ , and two expectations  $E_{\mathcal{D}}[f_i(\boldsymbol{\chi})]$  and  $E_{\boldsymbol{\theta}}[f_i]$ .

Both  $Z(\theta)$  and  $E_{\theta}[f_i]$  require inference to compute probabilities of each  $\xi$ 

### Exact Inference: Belief Propagation



#### 1. Gibbs Distribution

$$P(A,B,C.D) = \frac{1}{Z}\phi_1(A,B) \cdot \phi_2(B,C) \cdot \phi_3(C,D) \cdot \phi_4(D,A)$$
where
$$Z = \sum_{A,B,C,D} \phi_1(A,B) \cdot \phi_2(B,C) \cdot \phi_3(C,D) \cdot \phi_4(D,A)$$

Z=7,201,840

Sepset

 $\boxed{\tilde{P}_{_{\Phi}}\!\left(A,B,C,D\right)\!=\phi_{_{\!1}}\!\left(A,B\right)\!\phi_{_{\!2}}\!\left(B,C\right)\!\phi_{_{\!3}}\!\left(C,D\right)\!\phi_{_{\!4}}\!\left(D,A\right)}$ 

Unormalized Distribution

Cluster  $C_2$ 

A	ssig	nme		Unnormalized		
$a^0$	$b^0$	$c^0$	$d^0$	300000		
$a^0$	$b^0$	$c^0$	$d^1$	300000		
$a^0$	$b^0$	$c^1$	$d^0$	300000		
$a^0$	$b^0$	$c^1$	$d^1$	30		
$a^0$	$b^1$	$c^0$	$d^0$	500		
$a^0$	$b^1$	$c^0$	$d^1$	500		
$a^0$	$b^1$	$c^1$	$d^0$	5000000		
$a^0$	$b^1$	$c^1$	$d^1$	500		
$a^1$	$b^0$	$c^0$	$d^0$	100		
$a^1$	$b^0$	$c^0$	$d^1$	1000000		
$a^1$	$b^0$	$c^1$	$d^0$	100		
$a^1$	$b^{0}$	$c^1$	$d^1$	100		
$a^1$	$b^1$	$c^0$	$d^0$	10		
$a^1$	$b^1$	$c^0$	$d^1$	100000		
$a^1$	$b^1$	$c^1$	$d^0$	100000		
$a^1$	$b^1$	$c^1$	$d^1$	100000		

Clique/Cluster C<sub>1</sub>

1.A,B,D

 $\{B,D\}$  2. B,C,D

### 2. Clique Tree (triangulated):

#### **Initial Potentials:**

Each  $\psi$  has every factor involving its arguments

$$\begin{aligned} & \psi_1 \left( A, B, D \right) = \phi_1 \left( A, B \right) \phi_2 \left( B, C \right) \phi_3 \left( C, D \right) \phi_4 \left( D, A \right) \\ & \psi_2 \left( B, C, D \right) = \phi_1 \left( A, B \right) \phi_2 \left( B, C \right) \phi_3 \left( C, D \right) \phi_4 \left( D, A \right) \end{aligned}$$

### Computing Clique Beliefs ( $\beta_i$ ), Sepset Beliefs ( $\mu_{i,i}$ )

$$\begin{vmatrix} \beta_1 \Big( A, B, D \Big) = \tilde{P}_{\Phi} \Big( A, B, D \Big) = \sum_{C} \psi_1 \Big( A, B, D \Big) = \sum_{C} \phi_1 (A, B) \phi_2 (B, C) \phi_3 (C, D) \phi_4 (D, A) \\ \text{e.g.}, \quad \beta_1 (a^0, b^0, d^0) = 300,000 + 300,000 = 600,000 \end{aligned}$$

$$\begin{aligned} \mu_{1,2}(B,D) &= \sum_{C_1 - S_{1,2}} \beta_1 \Big( C_1 \Big) = \sum_A \beta_1 \Big( A, B, D \Big) \\ \text{e.g., } \mu_{1,2}(b^0, d^0) &= 600,000 + 200 = 600,200 \end{aligned}$$

$$\begin{split} \beta_2 \Big( B, C, D \Big) &= \tilde{P}_{\!_{\Phi}} \Big( B, C, D \Big) = \sum_{A} \mu_{1,2} \Big( B, D \Big) \cdot \psi_2 \Big( B, C, D \Big) = \sum_{A} \psi_2 \Big( B, C, D \Big) \\ e.g., \beta_2 \Big( b^0, c^0, d^0 \Big) &= 300,000 + 100 = 300,100 \end{split}$$

#### All Clique and Sepset Beliefs

Assignment	$\max_C$			Assignment	
$a^0   b^0   d^0  $	600,000			$b^0 \mid c^0 \mid d^0$	300.1
$  a^0   b^0   d^1  $	300,030	Assignment	$\max_{A,C}$	$ b^0  c^0  d^1 $	1.300.0
$  a^0   b^1   d^0  $	5,000,500	$b^0 \mid d^0$	600, 200	$ b^0  c^1  d^0 $	300.1
$  a^0   b^1   d^1  $	1,000	$b^0 \mid d^1$	1,300,130	$ b^0  c^1  d^1 $	
$  a^1   b^0   d^0  $	200	$b^1 d^0$	5, 100, 510	$ b^1 c^0 d^0$	
$  a^1   b^0   d^1  $	1,000,100	$b^1 d^1$	201,000	$b^{1} c^{0} d^{1}$	100.3
$  a^1   b^1   d^0  $	100,010			$b^{1} c^{1} d^{0}$	5, 100.
$\begin{vmatrix} a^1 & b^1 & d^1 \end{vmatrix}$	200,000			$b^1 \mid c^1 \mid d^1$	100.3
$\beta_1(A,B)$	(D)	$\mu_{1,2}(I$	$\beta_2(B,$	$\beta_2(B,C,\mathbf{D})$	

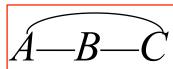
 $\tilde{P}_{\!\scriptscriptstyle \Phi}\!\left(\chi\right) \! = \! \frac{\prod\limits_{i \in V_T} \beta_i\!\left(C_i\right)}{\prod\limits_{(i-j) \in E_T} \mu_{i,j}\!\left(S_{i,j}\right)}$ 

Verifying Inference

$$\begin{split} &\tilde{P}_{\Phi}\left(a^{1},b^{0},c^{1},d^{0}\right) = 100 \\ &\frac{\beta_{1}\left(a^{1},b^{0},d^{0}\right)\beta_{2}\left(b^{0},c^{1},d^{0}\right)}{\mu_{1,2}\left(b^{0},d^{0}\right)} = \frac{200\cdot300\cdot100}{600\cdot200} = 100 \end{split}$$

# Example of Computing $E_{\mathcal{D}}[f_i(\chi)]$

 $\chi = \{A, B, C\}$ ; Pairwise Markov Network A - B



- Variables are binary
- Three clusters:  $C_1 = \{A, B\}, C_2 = \{B, C\}, C_3 = \{C, A\}$ 
  - Log-linear model  $P_{\theta}(\chi) = \frac{1}{Z(\theta)} \exp \left[ -\sum_{i=1}^{k} \theta_{i} f_{i}(D_{i}) \right]$  with two features:
  - $f_{00}(x,y)=1$  if x=0, y=0 and 0 otherwise and
  - $f_{11}(x,y)=1$  if x=1, y=1 and 0 otherwise

Both shared over all clusters

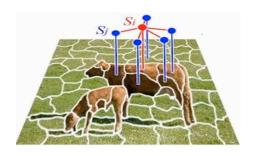
- Assume we have three data instances (A,B,C):
- $(0,0.0) \rightarrow$  Cluster AB, Cluster BC, Cluster CA have  $f_{00}(x,y)=1$
- (0,1,0) →Only Cluster CA has  $f_{00}(x,y)=1$
- (1,0,0) → Only Cluster BC has  $f_{00}(x,y)=1$ 
  - Unnormalized empirical feature counts, pooled over all clusters:  $E_{\tilde{z}} \lceil f_{zz} \rceil = (3+1+1)/3 = 5/3$  Can normalize using counts

$$\begin{aligned} E_{\tilde{P}} \left[ f_{00} \right] &= (3+1+1) / 3 = 5 / 3 \\ E_{\tilde{P}} \left[ f_{11} \right] &= (0+0+0) / 3 = 0 \end{aligned}$$

Can normalize using counts over all features i.e., (5/3)+0

### Difficulty with Exact Methods

- Exact Parameter Estimation Methods assume ability to compute
  - 1. Partition function  $Z(\theta)$  and
  - 2. Expectations  $E_{P_{\theta}}[f_i]$
- In many applications structure of network does not allow exact computation of these terms
  - In image segmentation, grid networks lead to exponential size clusters for exact inference



Cluster graph is clique tree with overlapping factors

### Discussion on Approximate Methods

### 1. In this section: approximate inference

- Decouple inference from Learning Parameters
  - Inference is a black-box
- But approximation may interfere with learning
  - Non-convergence of inference can lead to oscillating estimates of the gradient & no learning convergence

### 2. Next section: approximate objective function

- Whose optimization doesn't require much inference
- Approximately optimizing the likelihood function can be reformulated as exactly optimizing an approximate objective

## Approximate Inference

- Learning with Approximate Inference methods
  - 1. Belief Propagation
  - 2. MAP-based Learning

### Approximate Inference: Belief Propagation

- Popular Approach for Approximate Inference is Belief Propagation and its variants
  - An algorithm from this family would be used for inference in the model resulting from the learning procedure
- We should use the same inference algorithm that will be used for querying it
  - Model trained with same inference algorithm is better than model trained with exact inference!
- BP is run with each iteration of Gradient Ascent to compute expected feature count  $E_{P_{\theta}}[f_i]$  10

### Difficulty with Belief Propagation

- In principle, BP is run in each Gradient Ascent iteration
  - to compute  $E_{P_{\theta}}[f_i]$  used in gradient computation
    - Due to family preservation property each feature must be a subset of a cluster  $C_i$  in the cluster graph
      - Hence to compute  $E_{P_{\theta}}[f_i]$  we can compute BP marginals over  $C_i$
- But BP often does not converge
  - Marginals derived often oscillate
    - Final results depend on where we stop
  - As a result gradient computed is unstable
    - Hurt convergence properties of gradient descent
    - Even more severe with line search

### Convergent alternatives to Belief Propagation

### We describe three BP methods:

- 1. Pseudo- moment matching
  - Reformulate task of learning with approximate inference as optimizing an alternative objective
- 2. Maximum Entropy approximation (CAMEL)
  - General derivation that allows us to reformulate ML with BP as a unified optimization problem with an approximate objective
- 3. MAP-based Learning
  - Approximate expected feature counts with their counts in the single MAP assignment in current MN

## Pseudo-moment Matching

- Begin with analysis of fixed points in learning
  - Converged BP beliefs must satisfy  $E_{\beta_i(C_i)}[f_{C_i}] = E_D[f_i(C_i)]$ 
    - Or  $\beta_i \left( c_i^j \right) = \hat{P} \left( c_i^j \right)$ Convergent point is a set of beliefs that match the data
- Define for each sepset  $S_{i,j}$  between  $C_i$  and  $C_j$

$$\phi_i \leftarrow \frac{\beta_i}{\mu_{i,j}}$$

$$C_1: A,B,D$$

$$S_{12} = \{B,D\}$$

$$C_2: B,C,D$$

- We use the final set of potentials as the parameterization of the Markov Network
- Provides a closed-form solution for both inference and learning
  - Cannot be used with parameter regularization, nontable factors or CRFs

## **BP** and Maximum Entropy

- A more general derivation that allows us to reformulate maximum likelihood learning with belief propagation as a unified optimization problem with an approximate objective
- Opens door to use of better approximation algorithms
- Start with dual of maximum-likelihood problem

### Tractable Max Entropy

- Assume a cluster graph U consisting of
  - a set of clusters  $\{C_i\}$  connected by sepsets  $S_{ij}$ .
- Rather than optimize maximum-entropy
  - over the space of distributions Q we optimize

$$H_{Q}(\chi) \approx \sum_{C_{i} \in U} H_{\beta_{i}}(C_{i}) - \sum_{C_{i} - C_{j} \in S} H_{\mu_{i,j}}(S_{i,j})$$

This is exact for tree cluster graph but approx otherwise

Approx-Maximum-Entropy:

$$\begin{array}{ll} \mathbf{Find} \;\; Q(\; \pmb{\chi}\;) \\ \mathbf{Maximizing} \;\; \sum_{C_i \in U} H_{\beta_i} \Big( C_i \Big) - \sum_{C_i - C_j \in U} H_{\mu_{i,j}} \Big( S_{i,j} \Big) \end{array}$$

Subject to 
$$E_Q[f_i] = E_D[f_i], i=1,..., k$$
  $Q \in Local(U)$ 

Approximation is exact when cluster graph is a tree
 Method known as CAMEL (Constrained Approx Max Entropy Learning)

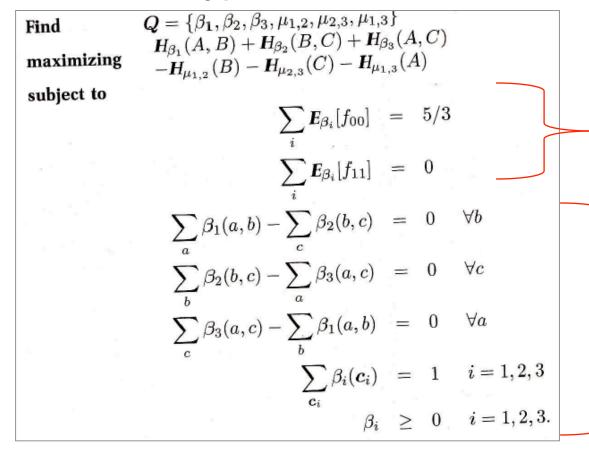
## Example of Max Ent Learning

- Pairwise Markov Network A—B—C
  - Variables are binary
  - Three clusters:  $C_1 = \{A,B\}, C_2 = \{B,C\}, C_3 = \{C,A\}$
  - Log-linear model with features
    - $f_{00}(x,y)=1$  if x=0, y=0 and 0 otherwise for x,y, instance of  $C_i$
    - $f_{11}(x,y)=1$  if x=1, y=1 and 0 otherwise
  - Three data instances (A,B,C): (0,0.0),(0,1,0),(1,0,0)
    - Unnormalized Feature counts, pooled over all clusters, are  $E_{\tilde{p}}[f_{00}] = (3+1+1)/3 = 5/3$

$$E_{\tilde{P}}[f_{11}] = (0+0+0)/3 = 0$$

## **CAMEL Optimization Problem**

- Optimization problem takes the form
  - with two types of constraints:



Type 1 Constraints:

$$E_{Q}[f_{i}]=E_{D}[f_{i}], i=1,..,k$$

Type 2 Constraints: Marginals from Cluster-graph approximation

### **CAMEL Solutions**

- CAMEL optimization is a constrained maximization problem with
  - linear constraints and
  - a nonconcave objective
- Several solution algorithms, one of which is
  - Lagrange multipliers for all constraints and optimize over resulting new variables

## Sampling-based Learning

We wish to maximize the log-likelihood

$$\ell(\theta:D) = \sum_{i} \theta_{i} \left( \sum_{m} f_{i} (\xi[m]) - M \ln Z(\theta) \right)$$

 In the sampling-based approach we use samples to estimate the partition function Z(θ) and use that estimate to perform gradient ascent

# Expressing $Z(\theta)$ as an expectation

- Partition function  $Z(\theta)$  is a summation over an exponentially large space
  - One approach to approximating this summation is to reformulate as expectation wrt a distribution  $Q(\chi)$

$$Z(\theta) = \sum_{\xi} \exp\left\{\sum_{i} \theta_{i} f_{i}(\xi)\right\}$$

$$= \sum_{\xi} \frac{Q(\xi)}{Q(\xi)} \exp\left\{\sum_{i} \theta_{i} f_{i}(\xi)\right\}$$

$$= E_{Q} \left[\frac{1}{Q(\chi)} \exp\left\{\sum_{i} \theta_{i} f_{i}(\chi)\right\}\right] \quad \text{Since } E_{Q} \left[g(\chi)\right] = \sum_{\xi} Q(\xi) g(\xi)$$

Since 
$$E_Q[g(\chi)] = \sum_{\xi} Q(\xi)g(\xi)$$

 This is precisely the form of importance sampling estimator 20

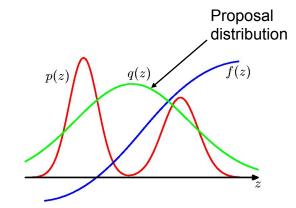
## Importance sampling

• To determine  $E_p[f] = \frac{1}{M} \sum_{m=1}^M f(\mathbf{z}^{[m]})$  from  $p(\mathbf{z})$ , we draw samples  $\{\mathbf{z}^{(m)}\}$  from a simpler dist.  $q(\mathbf{z})$ 

$$E[f] = \int f(z)p(z)dz$$

$$= \int f(z)\frac{p(z)}{q(z)}q(z)dz$$

$$= \frac{1}{M}\sum_{m=1}^{M}\frac{p(z^{(m)})}{q(z^{(m)})}f(z^{(m)})$$



Unlike rejection sampling All of the samples are retained

- Samples are weighted by ratios  $r_l = p(\mathbf{z}^{(l)}) \ / \ q(\mathbf{z}^{(l)})$ 
  - Known as importance weights
    - Which corrects the bias introduced by wrong distribution

## Importance Sampling Estimator

- We can approximate the partition function by generating samples from Q and correcting appropriately via weights
- We can simplify this expression by choosing Q to be  $P_{\theta^0}$  for some set of parameters  $\theta^0$

$$\begin{split} Z\!\left(\theta\right) &= E_{P_{\theta^0}}\!\left[ \frac{Z\!\left(\theta^0\right) \! \exp\left\{\! \sum_i \theta_i f_i\!\left(\chi\right)\!\right\}}{\exp\left\{\! \sum_i \theta_i^0 f_i\!\left(\chi\right)\!\right\}} \right] \\ &= \! Z\!\left(\theta^0\right) \! E_{P_{\theta^0}}\!\left[ \exp\left\{\! \sum_i \left(\theta_i - \theta_i^0\right) \! f_i\!\left(\chi\right)\!\right\} \right] \end{split}$$

# Samples to approximate $\ln Z(\theta)$

- If we can sample instances  $\xi^1,...\xi^K$  from  $P_{\theta^0}$ 
  - We can approximate the log-partition function as

$$\left|\ln Z\!\left(\theta\right) \approx \ln\!\left(\frac{1}{K}\!\sum_{k=1}^{K}\exp\left\{\!\sum_{i}\!\left(\theta_{i}-\theta_{i}^{0}\right)\!f_{i}\!\left(\xi^{k}\right)\!\right\}\!\right)\!+\ln Z\!\left(\theta^{0}\right)\right|$$

• We can plug this approximation of  $\ln Z(\theta)$  into the log-likelihood  $\frac{1}{M}\ell(\theta:D) = \sum_{i} \theta_{i} \left( E_{D}[f_{i}(d_{i})] \right) - \ln Z(\theta)$ 

and optimize it

• Note that  $\ln Z(\theta^0)$  is a constant

where  $E_D[f_i(\mathbf{d_i})]$  is the empirical expectation of f, i.e., its average in the data set

– that we can ignore in the optimization and the resulting expression is a simple function of  $\theta$  which can be optimized using gradient ascent

## Gradient ascent + Sampling

• Gradient ascent over  $\theta$  relative to

$$\left|\ln Z\!\left(\theta\right) \approx \ln\!\left(\frac{1}{K}\!\sum_{k=1}^{K}\exp\!\left\{\!\sum_{i}\!\left(\theta_{i}-\theta_{i}^{0}\right)\!f_{i}\!\left(\xi^{k}\right)\!\right\}\!\right)\!+\ln Z\!\left(\theta^{0}\right)\right|$$

 is equivalent to utilizing an importance sampling estimator directly to approximate the expected counts in the gradient of equation

$$\left| \frac{\partial}{\partial \theta_i} \frac{1}{M} \ell(\theta : D) = E_D[f_i(\chi)] - E_{\theta}[f_i] \right|$$

 However, it is generally it is useful to view such methods as exactly optimizing an approximate objective rather than approximately optimizing the exact likelihood

### Quality of importance sampling estimator

- It depends on the difference between  $\, heta \,$  and  $\, heta \,$   $\, heta \,$
- The greater the difference, the larger the variance of the importance weights
- Thus this type of approximation is reasonable only in a neighborhood of surrounding  $\theta^{\,0}$

## How to use this approximation?

- Iterate between two steps
  - 1. Use a sampling procedure to generate samples from current parameter set  $\theta^{t}$
  - 2. Then use gradient descent to find  $\theta^{t+1}$  that improves the approximate log-likelihood based on the samples
    - We can then regenerate samples and repeat the process.
  - As the samples are regenerated from a new distribution,
    - we can hope that that they are generated from a distribution not too far from the one we are currently optimizing maintaining a reasonable approximation

## MAP-based Learning

- Another approach to inference in learning
  - Approximating expected feature counts with the counts in the single MAP assignment to current MN
- Approximate gradient at assignment  $\theta$  is

$$E_D [f_i(\chi)] - f_i(\xi^{MAP}(\theta))$$

- where  $\xi^{MAP}(\theta)$  = arg max  $\xi P(\xi \mid \theta)$  is the MAP assignment given the current set of parameters  $\theta$
- Approach also called as Viterbi training
- Equivalent to exact optimization of approximate objective  $\frac{1}{M}\ell(\theta:D) - \ln P(\xi^{MAP}(\theta)|\theta)$

### Ex: CRFs for Protein Structure

- Predict the 3\_D structure of proteins
  - Proteins are chains of residues, each containing one of 20 possible amino acids
    - Amino acids are linked together into a common backbone structure onto which amino-specific side-chains are attached
  - Predict the side-chain conformations given the backbone.
  - Full configuration consists of upto four angles each of which takes on a continuous value
    - Discretize angle into a small no (3) rotamers
- Address optimization as MAP for CRF