Inference in Computer Science and Systems Biology Part I

Alfredo Braunstein

June 11, 2012

Outline



- Inference of Gene Regulation networks
- Inference of protein structure from protein families sequences
- 2 Bayesian inference
 - Bayes
 - Likelihood
 - Examples
 - Complex models
- 3 Approximate direct inference
 - Belief propagation
- 4 Inverse inference of trees
- 5 Coming up with models: maximum entropy principle
 - Observations
 - Examples
- 6 Other network reconstruction methods
- 7 Insufficient data

- Motivation

Inference of Gene Regulation networks

Transcriptional Gene Regulation

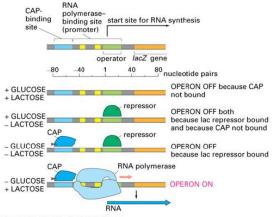


Figure 8-9 Essential Cell Biology, 2/e. (© 2004 Garland Science)

Inference in Computer Science and Systems Biology Part I Motivation Inference of Gene Regulation networks

Expression Data

- Identifying each precise regulation mechanism by experiments is very costly and time consuming: too many genes, way too many possible interactions!
- Hope to *infer* regulatory mechanisms from whole genome-scale experiments: *microarrays*

		172 stress conditions		
6152 genes	YAL001C	1.53	-0.06	
	YAL002W	-0.01	-0.30	
	YAL004W	0.24	0.76	•••
	:	:	:	

Yeast Dataset from: Grasch, Spellman, Mol. Biol. Cell (2000)

■ Log-ratios of expression data: overexpression, underexpression.

- Motivation

Inference of Gene Regulation networks

Inference of the gene-regulatory network

Two main goals:

- Inference of topology: Who regulates who?
- Inference of behaviour: predict the expression of a gene given the expression of other genes
- These are method of *inverse* inference: infer the model from the data

Inference of Gene Regulation networks

Inference of topology

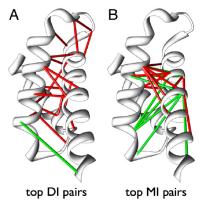
One way to do this is using coexpression networks.

- Compute the Pearson correlation coefficient *C_{ij}* for every pair *i*,*j* of genes
- Potential regulators of a gene are most correlated inputs
- Build the network of links for which $|C_{ij}|$ is above a certain treshold.
- But we can do better!

- Motivation

Inference of protein structure from protein families sequences

Inference of protein structure from protein families sequences



F. Morcos, A.Pagnani et al, 2011

Conditional probability

Conditional probability: restriction of a probability distribution to a subspace *B*:

$$P(A|B) = \frac{P(A \cap B)}{P(B)} = \frac{P(A,B)}{P(B)}$$

"Probability of A given B"

Example

What is the probability of the output of a die to be ≥ 2 given that the result is odd?

$$P(d \ge 2|d \text{ odd}) = P(d \ge 2, d \text{ odd}) / P(d \text{ odd}) = \frac{2}{6} / \frac{1}{2} = \frac{2}{3}$$

Bayes

$$P(A|B) = \frac{P(A,B)}{P(A)} = P(B|A)\frac{P(A)}{P(B)}$$

Example

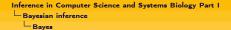
You are tested for an illness that is very rare (about 1:100000) with a fairly precise test (99% accuracy in both cases). You come up positive, yuck! Probability of illness? (a) 99% (b) 90% (c) 10% (d) 1% (e) 0.1%

$$P(I|+) = P(+|I)P(I)P(+)^{-1} \approx 0.99 \cdot 10^{-5} (0.01)^{-1} \approx 10^{-3}!$$

$$P(+) = P(+,I) + P(+, \text{not } I)$$

$$= P(+|I)P(I) + P(+| \text{not } I)P(\text{not } I)$$

$$= 0.99 \times 10^{-5} + 0.01 \times (1 - 10^{-5}) \approx 0.01$$



Bayes' rule in inference

- D = data, S = stochastic "machine", P(D|S) = stochastic rule, P(S) prior information about S
- A double stochastic process:
 - **1** S is extracted from P(S)
 - **2** D is extracted from P(D|S)

We observe only D. What can we guess about S?

$$\widetilde{P(S|D)} = \frac{P(D|S)P(S)}{P(D)} \propto \widetilde{P(D|S)P(S)}$$

Just the maths of common sense!

Bayes' rule iterated

Suppose we have the following multiple stochastic process:

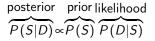
S is extracted from P(S)
D¹,...,D^M are extracted i.i.d from P(D|S)

$$P(S|D^1,\ldots,D^M) = \frac{P(D^1,\ldots,D^M|S)}{P(D)}P(S) \propto P(S)\prod_{\mu=1}^M P(D^\mu|S)$$

Sometimes it is written in *update* form:

$$P(S|D^{1},...,D^{M}) \propto P(D^{M}|S) \left(P(S)\prod_{\mu=1}^{M-1}P(D^{\mu}|S)\right)$$
$$= P(D^{M}|S) P(S|D^{1},...,D^{M-1})$$

MAP vs. Max likelihood



■ Maximum A Posteriori (MAP):

 $(\arg) \max_{S} P(S|D)$

Maximum Likelihood (ML):

 $(\arg) \max_{S} P(D|S)$

- ML=MAP for *uniform* prior, when it makes sense
- Two "Schools of thought"

Example: biased coins

I have two coins with head probabilities $p_1 = 0.5$ and $p_2 = 0.2$.

- 1 choose one at random with P(1) = 0.6, P(2) = 0.4.
- **2** I flip the coin and the output is tail.

Can we say something about the coin?

$$P(1|\text{tail}) \propto P(\text{tail}|1)P(1) = 0.5 \times 0.6 = 0.30$$

$$P(2|\text{tail}) \propto P(\text{tail}|2)P(2) = 0.8 \times 0.4 = 0.32$$

$$P(1|\text{tail}) = 0.30/(0.30 + 0.32) = 0.484$$

$$P(2|\text{tail}) = 0.32/(0.30 + 0.32) = 0.516$$

Not much!

Binomial distribution

Consider the outcome of n p-biased coins. The probability of k heads is

$$P(k|p) = \binom{n}{k} p^{k} (1-p)^{n-k}$$

Uniform prior

If P(p) = uniform, likelihood=posterior!

$$\mathcal{P}\left(k_{1},\ldots,k_{M}|p
ight) \propto p^{\sum_{\mu=1}^{M}k_{\mu}}\left(1-p
ight)^{\sum_{\mu=1}^{M}n-k_{\mu}} = \left(p^{\tilde{k}}\left(1-p
ight)^{n-\tilde{k}}
ight)^{M}$$

Binomial distribution

$$P(k_1,\ldots,k_M|p) \propto \left(p^{\tilde{k}} (1-p)^{n-\tilde{k}}\right)^M$$

with $\tilde{k} = \frac{1}{M} \sum_{\mu=1}^{M} k_{\mu}$ heads, the ML is attained at the max of

$$\mathscr{L} = \tilde{k} \log p + \left(n - \tilde{k}\right) \log \left(1 - p\right)$$

Let us find critical points:

$$0 = \frac{\partial \mathscr{L}}{\partial p} = \frac{\tilde{k}}{p} - \frac{\left(n - \tilde{k}\right)}{1 - p}$$

So $\frac{n - \tilde{k}}{\tilde{k}} = \frac{1 - p}{p}$, i.e. $p = \frac{\tilde{k}}{n}$.

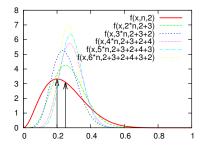
Note!

$$\langle p \rangle = \frac{\int_{0}^{1} p p^{M\tilde{k}} (1-p)^{M(n-\tilde{k})} dp}{\int_{0}^{1} p^{M\tilde{k}} (1-p)^{M(n-\tilde{k})} dp} = \frac{M\tilde{k}+1}{Mn+2}$$

Bayesian inference

Examples

Binomial



gnuplot code

```
f(p,n,k)=p**k*(1-p)**(n-k)/(k!*(n-k)!/(n+1)!)
pml(n,k)=k*1./n
pav(n,k)=(k+1)*1.0/(n+2)
n=10;k=2;
set arrow from pml(n,k),0 to pml(n,k), f(pml(n,k), n, k)
set arrow from pav(n,k),0 to pav(n,k), f(pav(n,k), n, k)
plot [0:1] f(x,n,2) lw 3, f(x,2*n,2+3), f(x,3*n,2+3+2),
f(x,4*n,2+3+2+4), f(x,5*n,2+3+2+4+3), f(x,6*n,2+3+2+4+3+2)
```

Example: Normal

$$P(x|(m,\sigma)) = \frac{1}{\sqrt{2\pi\sigma}} e^{-\frac{1}{2\sigma^2}(x-m)^2}$$

Given x^1, \ldots, x^M , we have

$$P(x^1,\ldots,x^M|(m,\sigma)) \propto e^{-\frac{1}{2\sigma^2}\sum_{\mu=1}^M (x^\mu-m)^2 - M\log\sigma}$$

If we try to maximize the log-likelihood $\mathscr{L}(m,\sigma) = -\frac{1}{2\sigma^2} \frac{1}{M} \sum_{\mu=1}^{M} (x^{\mu} - m)^2 - \log \sigma$

$$0 = \frac{\partial \mathscr{L}}{\partial m} = \frac{1}{\sigma^2} \frac{1}{M} \sum_{\mu=1}^{M} (x^{\mu} - m) \qquad 0 = \frac{\partial \mathscr{L}}{\partial \sigma} = \sigma^{-3} \left(\frac{1}{M} \sum_{\mu=1}^{M} (x^{\mu} - m)^2 - \sigma^2 \right)$$

I.e.
$$m = \frac{1}{M} \sum_{\mu}^{M} x^{\mu}, \sigma = \sqrt{\frac{1}{M} \sum_{\mu=1}^{M} (x^{\mu} - m)^2}$$

• What is the likelihood of (m, σ) when M = 1?

ML and KL divergence

Remember the KL divergence

$$KL(P||Q) = \sum_{\mathbf{x}} P(\mathbf{x}) \log \frac{P(\mathbf{x})}{Q(\mathbf{x})}$$

Assume you have a set of sample data \mathbf{x}^{μ} for $\mu = 1, ..., M$. Then consider the distribution $P(\mathbf{x}) = \frac{1}{M} \sum_{\mu=1}^{M} \delta(\mathbf{x}, \mathbf{x}^{\mu})$, and a distribution Q_{θ} parametrized by θ

$$\begin{split} \mathcal{K}\mathcal{L}(P||Q_{\theta}) &= \sum_{\mathbf{x}} \sum_{\mu=1}^{M} \delta\left(\mathbf{x}, \mathbf{x}^{\mu}\right) \log \frac{\sum_{\mu'=1}^{M} \delta\left(\mathbf{x}, \mathbf{x}^{\mu'}\right)}{Q_{\theta}\left(\mathbf{x}\right)} \\ &= -\log \prod_{\mu=1}^{M} Q_{\theta}\left(\mathbf{x}^{\mu}\right) \end{split}$$

That is, ML is the same as minimizing the KL divergence with $\frac{1}{M}\sum_{\mu=1}^{M}\delta\left(\mathbf{x},\mathbf{x}^{\mu}\right)!$

Ising model

Suppose given σ^1,\ldots,σ^M samples, and assume they were generated independently by an Ising model

$$P_{\boldsymbol{J},\boldsymbol{h}}(\boldsymbol{\sigma}) = Z_{\boldsymbol{J},\boldsymbol{h}}^{-1} e^{\sum_{i < j} J_{ij} \sigma_i \sigma_j + \sum_i h_i \sigma_i}$$

$$\begin{split} P\left(\sigma^{1},\ldots,\sigma^{M}|\boldsymbol{J},\boldsymbol{h}\right) &= \prod_{\mu=1}^{M} e^{\sum_{i < j} J_{ij}\sigma_{i}^{\mu}\sigma_{j}^{\mu}+\sum_{i}h_{i}\sigma_{i}^{\mu}-\log Z_{\boldsymbol{J},\boldsymbol{h}}} \\ &= e^{M\left(\sum_{i < j} J_{ij}\tilde{c}_{ij}+\sum_{i}h_{i}\tilde{m}_{i}-\log Z_{\boldsymbol{J},\boldsymbol{h}}\right)} \end{split}$$

- Depends only on the experimental first (\tilde{m}_i) and second moments (\tilde{c}_{ij}) of the data!
- The log-likelihood

$$\mathscr{L}(\boldsymbol{J},\boldsymbol{h}) = M\left(\sum_{i < j} J_{ij}\tilde{c}_{ij} + \sum_{i} h_i \tilde{m}_i - \log Z_{\boldsymbol{J},\boldsymbol{h}}\right)$$

How can we find **J**, **h** of maximum likelihood?

Ising Likelihood

$$\mathscr{L}(\boldsymbol{J}, \boldsymbol{h}) = M\left(\sum_{i < j} J_{ij}\tilde{c}_{ij} + \sum_{i} h_i \tilde{m}_i - \log Z_{\boldsymbol{J}, \boldsymbol{h}}\right)$$

Lets try to find critical points:

$$0 = \frac{\partial \mathscr{L}}{\partial J_{ij}} = M\left(\tilde{c}_{ij} - \frac{\partial \log Z_{J,h}}{\partial J_{ij}}\right) = M\left(\tilde{c}_{ij} - \langle \sigma_i \sigma_j \rangle\right) \qquad 0 = \frac{\partial \mathscr{L}}{\partial h_i} = M\left(\tilde{m}_i - \langle \sigma_i \rangle\right)$$

- Better: $-\log Z_{J,h}$ is a concave (\cap) function on J,h (and so is \mathcal{L}), so we can use gradient ascent!
- Unfortunately, estimating (σ_iσ_j) and (σ_i) is computationally hard! (NP-Complete). Possibilities:
 - **1** Exact enumeration (up to $N \approx 30$)
 - 2 Monte-Carlo methods (slow!)
 - 3 Mean-field type approximations (e.g. Belief Propagation)

Boltzmann learning

Boltzmann learning algorithm

- **1** (init) Set J = 0, h = 0
- (direct inference) **somehow** estimate $\{\langle \sigma_i \sigma_j \rangle\}_{i < j}$ and $\{\langle \sigma_i \rangle\}_i$ from $P_{J,h}$
- 4 (end?) if $|\Delta J_{ij}| < \varepsilon$ for all i < j, $|\Delta h_i| < \varepsilon$ for all i, exit
- **5** (update) $\mathbf{J} \leftarrow \mathbf{J} + \eta \Delta \mathbf{J}, \ \mathbf{h} \leftarrow \mathbf{h} + \eta \Delta \mathbf{h}$
- 6 Go to 2

But we need an (approximate) inference method for **2**!

Inference in Computer Science and Systems Biology Part I — Approximate direct inference

Belief propagation

Example: 3-Coloring (Potts)

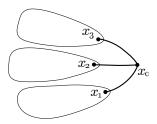
- Given a (finite) undirected graph G = (V, E)
- A proper 3-coloring is $\sigma_i \in \{\bullet, \bullet, \bullet\}$ for $i \in V$ such that $\sigma_i \neq \sigma_j$ if $(i,j) \in E$

$$P(\sigma) = \frac{1}{Z} \prod_{(ij) \in E} (1 - \delta(\sigma_i, \sigma_j))$$

- Hard computational problems (NP-Complete):
 - Finding a proper coloring
 - Estimating $P(\sigma_i, \sigma_j)$
 - Counting proper colorings
 - Deciding if there is at least one proper coloring!

Approximate direct inference

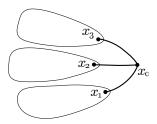
Belief propagation



$$\begin{split} N_{0}(\bullet) &= N^{(0)}(\bullet\bullet\bullet) + N^{(0)}(\bullet\bullet\bullet) + N^{(0)}(\bullet\bullet\bullet) + N^{(0)}(\bullet\bullet\bullet) + \cdots \\ &= N_{1}^{(0)}(\bullet) N_{2}^{(0)}(\bullet) N_{3}^{(0)}(\bullet) + N_{1}^{(0)}(\bullet) N_{2}^{(0)}(\bullet) N_{3}^{(0)}(\bullet) + \cdots \\ &= \left(N_{1}^{(0)}(\bullet) + N_{1}^{(0)}(\bullet)\right) \left(N_{2}^{(0)}(\bullet) + N_{2}^{(0)}(\bullet)\right) \left(N_{3}^{(0)}(\bullet) + N_{3}^{(0)}(\bullet)\right) \\ N_{0}(\bullet) &= \left(N_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet)\right) \left(N_{2}^{(0)}(\bullet) + N_{2}^{(0)}(\bullet)\right) \left(N_{3}^{(0)}(\bullet) + N_{3}^{(0)}(\bullet)\right) \\ N_{0}(\bullet) &= \left(N_{1}^{(0)}(\bullet) + N_{1}^{(0)}(\bullet)\right) \left(N_{2}^{(0)}(\bullet) + N_{2}^{(0)}(\bullet)\right) \left(N_{3}^{(0)}(\bullet) + N_{3}^{(0)}(\bullet)\right) \\ \end{split}$$

Approximate direct inference

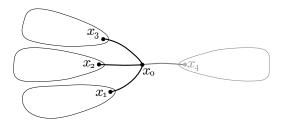
Belief propagation



$$\begin{split} P_{0}(\bullet) & \propto & P^{(0)}(\bullet\bullet\bullet) + P^{(0)}(\bullet\bullet\bullet) + P^{(0)}(\bullet\bullet\bullet) + P^{(0)}(\bullet\bullet\bullet) + \cdots \\ & = & P_{1}^{(0)}(\bullet) P_{2}^{(0)}(\bullet) P_{3}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) P_{2}^{(0)}(\bullet) P_{3}^{(0)}(\bullet) + \cdots \\ & = & \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) \right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet) \right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet) \right) \\ P_{0}(\bullet) & \propto & \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) \right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet) \right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet) \right) \\ P_{0}(\bullet) & \propto & \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) \right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet) \right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet) \right) \end{split}$$

- Approximate direct inference

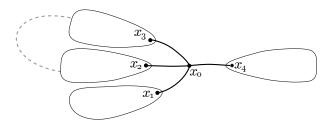
Belief propagation



$$\begin{split} P_{0}^{(4)}(\bullet) &\propto P^{(0)}(\bullet\bullet) + P^{(0)}(\bullet\bullet) + P^{(0)}(\bullet\bullet) + P^{(0)}(\bullet\bullet) + P^{(0)}(\bullet\bullet) + \cdots \\ &= P_{1}^{(0)}(\bullet) P_{2}^{(0)}(\bullet) P_{3}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) P_{2}^{(0)}(\bullet) P_{3}^{(0)}(\bullet) + \cdots \\ &= \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet)\right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet)\right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet)\right) \\ P_{0}^{(4)}(\bullet) &\propto \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet)\right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet)\right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet)\right) \\ P_{0}^{(4)}(\bullet) &\propto \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet)\right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet)\right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet)\right) \\ \end{split}$$

- Approximate direct inference

Belief propagation



$$\begin{split} P_{0}^{(4)}(\bullet) & \propto & P^{(0)}(\bullet\bullet\bullet) + P^{(0)}(\bullet\bullet\bullet) + P^{(0)}(\bullet\bullet\bullet) + P^{(0)}(\bullet\bullet\bullet) + \cdots \\ & \simeq & P_{1}^{(0)}(\bullet) P_{2}^{(0)}(\bullet) P_{3}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) P_{2}^{(0)}(\bullet) P_{3}^{(0)}(\bullet) + \cdots \\ & = & \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) \right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet) \right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet) \right) \\ P_{0}^{(4)}(\bullet) & \propto & \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) \right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet) \right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet) \right) \\ P_{0}^{(4)}(\bullet) & \propto & \left(P_{1}^{(0)}(\bullet) + P_{1}^{(0)}(\bullet) \right) \left(P_{2}^{(0)}(\bullet) + P_{2}^{(0)}(\bullet) \right) \left(P_{3}^{(0)}(\bullet) + P_{3}^{(0)}(\bullet) \right) \end{aligned}$$

Inference in Computer Science and Systems Biology Part I Approximate direct inference

Belief propagation

BP Equations (coloring)

$$q_{ij}(\sigma_i) \propto \psi_i(\sigma_i) \prod_{k \in \partial i \setminus j} \sum_{\sigma_k \neq \sigma_i} q_{ki}(\sigma_k)$$

This system is a fixed point $\mathbf{F}(\mathbf{q}) = \mathbf{q}$ equation for $\mathbf{q} = \{q_{ij}, q_{ji}\}_{(ij)\in E} \in [0, 1]^{2|E|}$ and is solved normally by iteration:

$$\mathbf{q}_{\infty} = \lim_{k o \infty} \mathbf{F}^{(k)} \left(\mathbf{q}_0
ight)$$

On a fixed point, we can compute

$$p_{i}(\sigma_{i}) \propto \Psi_{i}(\sigma_{i}) \prod_{k \in \partial i} \sum_{\sigma_{k} \neq \sigma_{i}} q_{ki}(\sigma_{k})$$
$$p_{ij}(\sigma_{i},\sigma_{j}) \propto q_{ij}(\sigma_{i}) q_{ji}(\sigma_{j}) (1 - \delta(\sigma_{i},\sigma_{j}))$$

Inference in Computer Science and Systems Biology Part I Approximate direct inference

- h C

Belief propagation

Belief Propagation (pairwise models)

Given a distribution:

$$P(\sigma) = \frac{1}{Z} \prod_{(ij)\in E} \psi_{ij}(\sigma_i, \sigma_j) \prod_i \psi_i(\sigma_i) = \frac{1}{Z} e^{-\left(\sum_{(ij)\in E} -\log \psi_{ij}(\sigma_i, \sigma_j) + \sum_i -\log \psi_i(\sigma_i)\right)}$$

BP Equations, pairwise potentials

$$\begin{array}{lll} q_{ij}(\sigma_i) & \propto & \psi_i(\sigma_i) \prod_{k \in \partial_i \setminus j} \sum_{\sigma_k} q_{ki}(\sigma_k) \, \psi_{ki}(\sigma_k, \sigma_i) \ (\text{message}) \\ p_i(\sigma_i) & \propto & \psi_i(\sigma_i) \prod_{k \in \partial_i} \sum_{\sigma_k} q_{ki}(\sigma_k) \, \psi_{ki}(\sigma_k, \sigma_i) \ (\text{marginal}) \\ p_{ij}(\sigma_i, \sigma_j) & \propto & \psi_{ij}(\sigma_i, \sigma_j) \, q_{ij}(\sigma_i) \, q_{ji}(\sigma_j) \ (\text{marginal}) \end{array}$$

Approximate direct inference

Belief propagation

BP for crosswords

- English dictionary D (set of english words)
- Indices: a set X of letters coordinates, one for each non-black square, a set H of horizontal words *indices*, one for each horizontal blank sequence, a set V of vertical word *indices*, one for each vertical blank sequence,
- **Variables**: $h_s \in D$ for each $s \in H$, $v_t \in D$ for each $t \in V$, $x_{ij} \in \{a, \ldots, z\}$ for each $ij \in X$
- For each non-black square *ij*,
 - $s(ij) \in H$ =crossing horizontal word, p(ij)= position of ij within,
 - $t(ij) \in V$ =crossing vertical word, q(ij) position of ij within
- Constraints: For each non black position ij: the following two conditions have to be ensured: (h_{s(ij)})_{p(ii)} = x_{ij} and (v_{t(ij)})_{g(ii)} = x_{ij}
- In summary: |H| + |V| + |X| variable nodes, 2|X| constraints

$$P(\mathbf{h}, \mathbf{v}, \mathbf{x}) = \frac{1}{Z} \prod_{ij \in X} \delta\left(\left(h_{s(ij)} \right)_{p(ij)}; x_{ij} \right) \delta\left(\left(v_{t(ij)} \right)_{q(ij)}; x_{ij} \right)$$

Inference in Computer Science and Systems Biology Part I — Approximate direct inference

Belief propagation

Exact inference on trees

Let T = (V, E) be a tree, and assume P a *T*-factorized distribution, i.e. $P(\sigma) = \frac{1}{Z} \prod_{(ij) \in E} \psi_{ij}(\sigma_i, \sigma_j)$. Then:

$$P(\sigma) = \prod_{(ij)\in E} \frac{P(\sigma_i, \sigma_j)}{P(\sigma_i) P(\sigma_j)} \prod_i P(\sigma_i)$$

For a general graph G, it is only an approximation!

■ It is called the *Bethe* approximation.

Approximate direct inference

Belief propagation

Entropy of a tree distribution

If P is T-factorized, then

$$-S(P) = \sum_{\sigma} P(\sigma) \ln P(\sigma)$$

=
$$\sum_{(ij)\in E} KL(P(\sigma_i, \sigma_j) || P(\sigma_i) P(\sigma_j)) - \sum_i S(P(\sigma_i))$$

=
$$\sum_{(ij)\in E} M_{ij} - \sum_i H_i$$

Approximate direct inference

Belief propagation

Average Energy and Free Energy

For every G = (V, E)-factorized Ising model,

$$\begin{array}{lll} -\langle H\rangle &=& \displaystyle\sum_{(ij)\in E} J_{ij} \langle \sigma_i \sigma_j \rangle + \displaystyle\sum_i h_i \langle \sigma_i \rangle \\ -\log Z_{J,h} &=& \langle H \rangle - S \\ &=& \langle H \rangle + \displaystyle\sum_{\sigma} P(\sigma) \log P(\sigma) \end{array}$$

If P is T-factorized, then

$$-\log Z_{J,h} = \langle H \rangle + \sum_{(ij) \in E} M_{ij} - \sum_{i} H_{i}$$

These expressions for S and $\log Z$ are exact for trees, just approximations for general graphs!

Inference in Computer Science and Systems Biology Part I
Approximate direct inference
Belief propagation

Mutual Information

Mutual Information is a measure of correlation:

$$MI(x,y) = \sum_{x} P(x,y) \log \frac{P(x,y)}{P(x)P(y)}$$

In terms of the KL divergence:

$$MI(x,y) = KL(P(x,y)||P(x)P(y))$$

It can be also thought as "information gain": how much information about x is gained (in average) by knowing the value of y:

$$MI(x,y) = S(P(x)) - \sum_{y} P(y)S(P(x|y)) \\ = S(P(y)) - \sum_{x} P(x)S(P(y|x))$$

 $MI(x,y) \le S(P(x))$ If x = y (i.e. $P(x,y) = \delta(x,y)P(x)$), MI(x,y) = S(P(x)). If P(x,y) = P(x)P(y), MI(x,y) = 0

Inference of trees

Suppose that we are told that some tree-factorized Ising model produced a set of samples:

$$\sigma^{1},\ldots,\sigma^{M}\sim P(\sigma)=rac{1}{Z_{\mathbf{J},\mathbf{h}}}e^{\sum_{i< j}J_{ij}\sigma_{i}\sigma_{j}+\sum_{i}h_{i}\sigma_{i}}$$

■ How do we find the tree T = (V, E) and the *T*-factorized \mathbf{J}, \mathbf{h} (i.e. such that $J_{ij} \neq 0 \implies (ij) \in E$) of ML?

Likelihood of a tree

Given samples $\sigma^1, \ldots, \sigma^M$, consider J^*, h^* the T = (V, E)-factorized ML couplings (T tree), then

$$\begin{aligned} \mathscr{L}(\boldsymbol{J}^*, \boldsymbol{h}^*) &= \sum_{(ij)\in \boldsymbol{E}} J_{ij}^* \tilde{c}_{ij} + \sum_i h_i^* \tilde{m}_i - \log Z_{\boldsymbol{J}^*, \boldsymbol{h}^*} \\ &= -\langle \tilde{H^*} \rangle - \log Z_{\boldsymbol{J}^*, \boldsymbol{h}^*} \\ &= -\langle \tilde{H^*} \rangle + \langle H^* \rangle - S^* \\ &= -\langle \tilde{H^*} \rangle + \langle H^* \rangle + \sum_{(ij)\in \boldsymbol{E}} M_{ij}^* + \sum_i S_i^* \end{aligned}$$

Chow-Liu (1968)

$$\mathscr{L}(\boldsymbol{J}^*, \boldsymbol{h}^*) = -\langle \tilde{H}^* \rangle + \langle H^* \rangle + \sum_{(ij) \in E} M^*_{ij} + \sum_i S^*_i$$

Two key observations:

■ We have seen that $\mathbf{P}_{\mathbf{J}^*,\mathbf{h}^*}$ must reproduce the first (\tilde{m}_i) and second (\tilde{c}_{ij}) moments of the data over T (so $\langle \tilde{H}^* \rangle = \langle H^* \rangle$). Then it must reproduce also $\tilde{P}(\sigma_i, \sigma_j) = \frac{1}{4} (\tilde{c}_{ij}\sigma_i\sigma_j + \tilde{m}_i\sigma_i + \tilde{m}_j\sigma_j + 1)$. In particular, $M_{ij}^* = \tilde{M}_{ij}$ and $S_i^* = \tilde{S}_i$.

2 The term \tilde{S}_i does not depend on T

$$\mathscr{L}(\boldsymbol{J}^*, \boldsymbol{h}^*) = \sum_{(ij)\in \boldsymbol{E}} \tilde{M}_{ij} + \text{const.}$$

And we want to maximize with respect to T (topology)

Maximum Spanning Tree (Kruskal 1956)

Given a connected graph G = (V, E) and weights $M : E \to \mathbb{R}_+$, finding the maximum spanning tree can be done as follows:

Kruskal's algorithm

Order edges so as to have $M_{e_1} \ge M_{e_2} \ge \cdots M_{e_{|E|}}$ Set $E' \leftarrow \emptyset$ For $s = 1, \dots, |E|$:
If $(V, E' \cup \{e_s\})$ has no loop: $E' \leftarrow E' \cup \{e_s\}$

At the end, (V,E') is a maximum spanning tree, i.e. a tree that maximizes $\sum_{e\in E'}M_e$

Putting all the bits toghether

1 Compute
$$M_{ij}$$
 for $i < j$

1 Use Kruskal to compute T the MST for the M_{ij}

$$\tilde{P}(\sigma_i,\sigma_j) = e^{J_{ij}\sigma_i\sigma_j + a_{ij}\sigma_i + b_{ij}\sigma_j + f_{ij}} \quad \tilde{P}(\sigma_i) = e^{h'_i\sigma_i + f_i}$$

$$P(\sigma) = \prod_{(ij)\in T} \tilde{P}(\sigma_i, \sigma_j) \prod_i \tilde{P}(\sigma_i)^{1-d_i}$$

$$\propto e^{\sum_{(ij)\in T} J_{ij}\sigma_i\sigma_j + a_{ij}\sigma_i + b_{ij}\sigma_j + \sum_i h'_i\sigma_i(1-d_i)}$$

$$\propto e^{\sum_{(ij)\in T} J_{ij}\sigma_i\sigma_j + \sum_i \sigma_i h_i}$$

where h_i is computed by collecting all coefficients of σ_i .

Maximum Spanning Tree

Proof by induction on $t: E' \subseteq E''$ for some MST E'' in every step t of Kruskal (assume that for some step $t \ge 0$, E' is included in an MST E'' and prove that $E' \cup \{e_t\}$ is also included in some MST)

- **I** If $E' \cup \{e_t\}$ is also included in E'', **done**. Otherwise:
- 2 $E'' \cup \{e_t\}$ has a loop p(E'' is a tree).
- **3** Take any edge f in $p \setminus (E' \cup \{e_t\})$ (such an edge must exist, otherwise $p \subseteq E' \cup \{e_t\}$).
- 4 We have $M_{e_t} \ge M_f$ (otherwise f would have been added before e_t).
- **5** $E''' = E'' \setminus \{f\} \cup \{e_t\}$ is a tree, $\sum_{(ij) \in E'''} M_{ij} \ge \sum_{(ij) \in E''} M_{ij}$, so E'''MST, and $E' \cup \{e_t\} \subseteq E'''$ done

Example

N = 5, M = 6, Data:

σ_1	σ_2	σ_3	σ_4	σ_5
1	1	1	$^{-1}$	1
1	-1	1	$^{-1}$	1
1	1	1	$^{-1}$	1
$^{-1}$	-1	1	1	1
$^{-1}$	1	$^{-1}$	1	$^{-1}$
$^{-1}$	1	1	$^{-1}$	$^{-1}$

Marginals:

$$\begin{split} P_{1} &= \frac{1}{6} \left(\begin{array}{c} 3\\ 3 \end{array}\right), P_{2} = \frac{1}{6} \left(\begin{array}{c} 2\\ 4 \end{array}\right), P_{3} = \frac{1}{6} \left(\begin{array}{c} 1\\ 5 \end{array}\right), P_{4} = \frac{1}{6} \left(\begin{array}{c} 4\\ 2 \end{array}\right), P_{5} = \frac{1}{6} \left(\begin{array}{c} 3\\ 3 \end{array}\right) \text{ and} \\ P_{12} &= \frac{1}{6} \left(\begin{array}{c} 1 & 2\\ 1 & 2 \end{array}\right), P_{13} = \frac{1}{6} \left(\begin{array}{c} 1 & 2\\ 0 & 3 \end{array}\right), P_{14} = \frac{1}{6} \left(\begin{array}{c} 1 & 2\\ 3 & 0 \end{array}\right), P_{15} = \\ \frac{1}{6} \left(\begin{array}{c} 2 & 1\\ 0 & 3 \end{array}\right), P_{23} = \frac{1}{6} \left(\begin{array}{c} 0 & 2\\ 1 & 3 \end{array}\right), P_{24} = \frac{1}{6} \left(\begin{array}{c} 1 & 1\\ 3 & 1 \end{array}\right), P_{25} = \frac{1}{6} \left(\begin{array}{c} 0 & 2\\ 2 & 2 \end{array}\right), P_{34} = \\ \frac{1}{6} \left(\begin{array}{c} 0 & 1\\ 4 & 1 \end{array}\right), P_{35} = \frac{1}{6} \left(\begin{array}{c} 1 & 0\\ 1 & 4 \end{array}\right), P_{45} = \frac{1}{6} \left(\begin{array}{c} 1 & 3\\ 1 & 1 \end{array}\right) \\ \text{Mutual information:} \ M_{15} = 0.459, M_{45} = 0.459, M_{35} = 0.317, M_{34} = 0.317, M_{25} = \\ 0.252, M_{13} = 0.191, M_{23} = 0.109, M_{45} = 0.044, M_{24} = 0.044, M_{12} = 0 \\ \text{Kruskal edges:} \ (15), (45), (35), (34), (25) \end{split}$$

Independent pairs

Assume the Bethe expression for trees to be valid for the complete graph:

$$P(\sigma|\boldsymbol{J},\boldsymbol{h}) = \prod_{i < j} \frac{P(\sigma_i, \sigma_j)}{P(\sigma_i) P(\sigma_i)} \prod_i P(\sigma_i)$$

we parametrize

$$P(\sigma_i, \sigma_j) = e^{J'_{ij}\sigma_i\sigma_j + a_{ij}\sigma_i + b_{ij}\sigma_j + f_{ij}} \quad P(\sigma_i) = e^{h''_i\sigma_i + f_i}$$

But then,

$$P(\boldsymbol{\sigma}|\boldsymbol{J},\boldsymbol{h}) = e^{\sum_{i < j} J'_{ij} \sigma_i \sigma_j + \sum_i (1 - d_i) \left(\sum_{j > i} \left(a_{ij} + b_{ji}\right) + h''_i\right) \sigma_i} \implies J'_{ij} = J_{ij}$$

but we know that on the point of ML, $P(\sigma_i \sigma_j) = \tilde{P}(\sigma_i, \sigma_j)$ so we can get J_{ij} directly from the data as in the two-spin system:

$$J_{ij} = \log rac{ ilde{p}_{++} ilde{p}_{--}}{ ilde{p}_{+-} ilde{p}_{-+}}$$

This exactly the same as if we consider each link separately (a single link is a tree!). This is called the **independent pairs** approximation.

BP on the Ising model

With the change of variables

$$h_{ij}=rac{1}{2}\lograc{q_{ij}\left(+1
ight)}{q_{ij}\left(-1
ight)}$$

The BP equations for the Ising model

$$q_{ij}(\sigma_i) \propto e^{h_i \sigma_i} \prod_{k \in \partial_i \setminus j} \sum_{\sigma_k} q_{ki}(\sigma_k) e^{J_{ki} \sigma_k \sigma_i}$$

become:

$$h_{ij} = h_i + \sum_{l \in \partial i \setminus j} \tanh^{-1} (\tanh J_{li} \tanh h_{li})$$
$$m_i = \tanh \left(h_i + \sum_{l \in \partial i} \tanh^{-1} (\tanh J_{li} \tanh h_{li}) \right)$$

Susceptibility Propagation

If we define

$$g_{ijk} = \frac{\partial h_{ij}}{\partial h_k}$$

Taking derivatives of the BP equations we obtain Susceptibility Propagation Equations (Mézard & Mora 2007):

$$g_{ijk} = \delta_{ik} + \sum_{l \in \partial_i \setminus j} g_{lik} \tanh J_{li} \frac{1 - \tanh^2 h_{li}}{1 - \tanh^2 J_{li} \tanh^2 h_{li}}$$

This gives a much better approximation for the susceptibility $\chi_{ij} = c_{ij} - m_i m_j = \frac{\partial m_i}{\partial h_j}:$ $\chi_{ij} = \left(\frac{\tanh J_{ij} + \tanh h_{ij} \tanh h_{ji}}{1 + \tanh J_{ii} \tanh h_{ii}} - m_i m_j\right) g_{jij} + g_{ijj} \left(1 - m_i^2\right)$

that can be employed for gradient ascent or on a coordinated $h_{ij}, g_{ijk}, J_{ij}, h_i$ updating scheme.

Inference in Computer Science and Systems Biology Part I Coming up with models: maximum entropy principle

Example

We will deal with partial observation of extractions from a distribution over $X = \{1, ..., n\}$.

- Suppose you see that over M samples, n_3 samples were the number 3. In the remaining $M n_3$, you just don't know.
- You need to point out **one** plausible distribution for the data.
- Would your guess be e.g. $P(k) = \frac{n_3}{M} \delta(k,3) + \frac{M-n_3}{M} \delta(k,2)$? This one is compatible with the observations!
- Or would you rather guess $P(k) = \frac{n_3}{M} \delta(k,3) + \frac{M-n_3}{M} (1 \delta(k,3))$, i.e. completely flat in the unobserved part?

Inference in Computer Science and Systems Biology Part I Coming up with models: maximum entropy principle

- Observations

Another example

Same setup as before.

- Suppose you only observe that over M samples, n_{23} samples were either 2 or 3, and n_{34} samples were either 3 or 4.
- How do we find the *flattest* possible distribution given the observations?

Coming up with models: maximum entropy principle

- Observations

General case: making predictions from partial observations

- How can we come up with reasonable models?
- Suppose we have a distribution $\mathbf{P} : \mathbf{X} \to [0,1]$ and we are given an observable for a variable f:

$$\bar{f} = \sum_{\mathbf{x}} f(\mathbf{x}) P(\mathbf{x})$$

How does one compute

$$\bar{g} = \sum_{\mathbf{x}} g(\mathbf{x}) P(\mathbf{x})?$$

But!

(very) undertermined system ($|\mathbf{X}|$ unknows, 2 equations)!

Coming up with models: maximum entropy principle

- Observations

Maximum Entropy

- Let us find the distribution **P** that satisfies $\bar{f} = \sum_{\mathbf{x}} f(\mathbf{x}) P(\mathbf{x})$ and $S(\mathbf{P}) = -\sum_{\mathbf{x}} P(\mathbf{x}) \ln P(\mathbf{x})$ is maximum (Jaynes 1957)
- This is the "less constrained / flattest distribution" compatible with the observation
- Using Lagrange multipliers...

$$\Gamma(\lambda,\mu,\mathbf{P}) = S(\mathbf{P}) + \mu\left(\overline{f} - \sum_{\mathbf{x}} f(\mathbf{x}) P(\mathbf{x})\right) + \lambda\left(1 - \sum_{\mathbf{x}} P(\mathbf{x})\right)$$

 And we need to find an unconstrained maximum for max_{λ,μ,P} Γ(λ,μ, P). Taking derivative w.r.t P(x)

$$0 = \frac{\partial \Gamma}{\partial P(\mathbf{x})} = -\ln P(\mathbf{x}) - P(\mathbf{x}) / P(\mathbf{x}) - \mu f(\mathbf{x}) - \lambda$$

$$P(\mathbf{x}) = e^{-\mu f(\mathbf{x}) - (1+\lambda)} \propto e^{-\mu f(\mathbf{x})}$$

• (A Boltzmann / exponential distribution!)

Inference in Computer Science and Systems Biology Part I Coming up with models: maximum entropy principle

Many observations

In general for many simultaneous observations f_1, \ldots, f_m ,

$$\max_{\lambda,\mu_{1},\dots,\mu_{m},\mathbf{P}} S(\mathbf{P}) + \sum_{a=1}^{m} \mu_{a} \left(\bar{f}_{a} - \sum_{\mathbf{x}} f_{a}(\mathbf{x}) P(\mathbf{x}) \right) + \lambda \left(1 - \sum_{\mathbf{x}} P(\mathbf{x}) \right)$$
$$0 = \frac{\partial \Gamma}{\partial P(\mathbf{x})} = -\ln P(\mathbf{x}) - P(\mathbf{x}) / P(\mathbf{x}) - \sum_{a=1}^{m} \mu_{a} f_{a}(\mathbf{x}) - \lambda, \text{ so}$$
$$P(\mathbf{x}) \propto e^{-\sum_{a=1}^{m} \mu_{a} f_{a}(\mathbf{x})}$$

Fact!

$$\sum_{\mathbf{x}} P(\mathbf{x}) \log \frac{P(\mathbf{x})}{\alpha} = -S - \log \alpha, \text{ i.e. } \min KL(P, \text{uniform}) = \max S$$

Coming up with models: maximum entropy principle

Examples

Going back to our n_{23} , n_{34} example

- Our observables were $f_1(i) = \delta(i,2) + \delta(i,3)$ and $f_2(i) = \delta(i,3) + \delta(i,4)$, and $\langle f_1 \rangle = \frac{n_{23}}{M} = p_{23}, \langle f_2 \rangle = \frac{n_{34}}{M} = p_{34}$
- Maximum entropy says:

$$P(i) \propto e^{-\mu_1(\delta(i,2) + \delta(i,3)) - \mu_2(\delta(i,3) + \delta(i,4))}$$

• i.e, defining
$$r = e^{-\mu_1}$$
 and $s = e^{-\mu_2}$ we get $P(i) = \frac{1}{Z} r^{\delta(i,2) + \delta(i,3)} s^{\delta(i,4) + \delta(i,3)}$

$$Z = \sum_{i=1}^{n} r^{\delta(i,2)+\delta(i,3)} s^{\delta(i,4)+\delta(i,3)} = r + rs + s + (n-3)$$

$$p_{23} = \frac{1}{Z} \sum_{i=1}^{n} (\delta(i,2) + \delta(i,3)) r^{\delta(i,2)+\delta(i,3)} s^{\delta(i,4)+\delta(i,3)} = \frac{1}{Z} (r+rs)$$

$$p_{34} = \frac{1}{Z} \sum_{i=1}^{n} (\delta(i,3) + \delta(i,4)) r^{\delta(i,2)+\delta(i,3)} s^{\delta(i,4)+\delta(i,3)} = \frac{1}{Z} (rs+s)$$

a 3×3 system (solve it!)

Coming up with models: maximum entropy principle

Examples

So

Example: ME distribution on \mathbb{N}_0 with fixed mean

Let P be the distribution of maximum entropy on $\{0, 1, ...\}$ with mean $m \ge 0$ (that is $m = \sum_i iP(i)$).

$$P(i) = \frac{1}{Z}e^{-\mu i} = \frac{1}{Z}(e^{-\mu})^{i}$$

Denote $r = e^{-\mu}$.

$$1 = \sum_{i=0}^{\infty} P(i) = \frac{1}{Z} \sum_{i=0}^{\infty} r^i$$

So $Z = \frac{1}{1-r}$, i.e. $P(i) = r^i (1-r)$. This is called the

So $Z = \frac{1}{1-r}$, i.e. $P(i) = r^i (1-r)$. This is called the *geometric* distribution.

$$m = \sum_{i=0}^{\infty} iP(i) = (1-r) r \sum_{i=0}^{\infty} ir^{i-1} = (1-r) r \frac{\partial}{\partial r} \left(\sum_{i=0}^{\infty} r^{i} \right)$$
$$= (1-r) r \frac{1}{(1-r)^{2}} = \frac{r}{1-r} = \frac{1}{1-r} - 1$$
$$r = 1 - \frac{1}{m+1}.$$

Inference in Computer Science and Systems Biology Part I
Coming up with models: maximum entropy principle
Examples

Example: spins, first moments

Suppose $\sigma_i \in \{-1,1\}$ for i = 1, ..., N, and we are given the N observables $m_i = \langle \sigma_i \rangle$ for i = 1, ..., N. Then the maximum entropy distribution is

$$P(\sigma) \propto e^{-\sum_i \mu_i \sigma_i} = \prod_i e^{-\mu_i \sigma_i}$$

As $m_i = \sum_{\sigma} P(\sigma) \sigma_i$,

$$m_{i} = \frac{\sum_{\sigma} \sigma_{i} \prod_{j} e^{-\mu_{j}\sigma_{j}}}{\sum_{\sigma} \prod_{j} e^{-\mu_{j}\sigma_{j}}}$$

$$= \frac{\sum_{\sigma^{-i}} \prod_{j \neq i} e^{-\mu_{j}\sigma_{j}} \sum_{\sigma_{i}} \sigma_{i} e^{-\mu_{i}\sigma_{i}}}{\sum_{\sigma^{-i}} \prod_{j \neq i} e^{-\mu_{j}\sigma_{j}} \sum_{\sigma_{i}} e^{-\mu_{i}\sigma_{i}}}$$

$$= \frac{\sum_{\sigma_{i}} \sigma_{i} e^{-\mu_{i}\sigma_{i}}}{\sum_{\sigma_{i}} e^{-\mu_{i}\sigma_{i}}} = \tanh(-\mu_{i})$$

So $\mu_i = -\tanh^{-1}(m_i)$.

Coming up with models: maximum entropy principle

Examples

Example: spins, first two moments

Suppose $\sigma_i \in \{-1,1\}$ for i = 1, ..., N, and we are given the $\frac{1}{2}N(N-1) + N$ observables $c_{ij} = \langle \sigma_i \sigma_j \rangle$ for $1 \le i < j \le N$ and $m_i = \langle \sigma_i \rangle$ for i = 1, ..., N. Then the maximum entropy distribution is

$$P(\sigma) \propto e^{\sum_{i < j} J_{ij}\sigma_i\sigma_j + \sum_i h_i\sigma_i}$$

i.e. an Ising model! This model has further restrictions on couplings and fields: $\sum_{\sigma} P(\sigma) \sigma_i \sigma_j = c_{ij}, \sum_{\sigma} P(\sigma) \sigma_i = m_i$

• We know how to find J_{ij} and h_i in the case of a tree prior...

Inference in Computer Science and Systems Biology Part I Coming up with models: maximum entropy principle Examples

Maximum Likelihood and Maximum Entropy

For an Ising model, we have seen that

$$\mathscr{L}(\mathsf{J},\mathsf{h}) = \sum_{i < j} \tilde{c}_{ij} J_{ij} + \sum_{i} \tilde{m}_{i} h_{i} - \log Z_{\mathsf{J},\mathsf{h}}$$

But also that on the point of ML, $\tilde{c}_{ij} = c^*_{ij} = \langle \sigma_i \sigma_j \rangle$ and $\tilde{m}_i = m^*_i = \langle \sigma_i \rangle$. So

$$\mathscr{L}(\mathbf{J}^*,\mathbf{h}^*) = -\langle E \rangle_{\mathbf{J}^*,\mathbf{h}^*} - \log Z_{\mathbf{J}^*,\mathbf{h}^*} = S(P_{\mathbf{J}^*,\mathbf{h}^*})$$

ML=ME

The **J**, **h** of ML describe the distribution of ME that reproduce $\tilde{m}_i, \tilde{c}_{ij}$

Inference in Computer Science and Systems Biology Part I — Other network reconstruction methods

ARACNE

Data Processing inequality: If P(x, y|z) = P(x|z)P(y|z) then $M_{xy} \le \min \{M_{xz}, M_{yz}\}$

- This can be used for reconstruction (Califano & al, 2006): for every triplet *i*,*j*,*k* consider *M*_{*ij*}, *M*_{*ik*}, *M*_{*jk*} and eliminate the smallest one.
- The resulting graph contains the Chow-Liu tree.
- Running time $\sim N^3$

Inference in Computer Science and Systems Biology Part I — Other network reconstruction methods

Reconstruction using independence

Observation

If $j \notin \partial i \cup \{i\}$

$$P(x_i, x_j | \mathbf{x}_{\partial i}) = P(x_i | \mathbf{x}_{\partial i}) P(x_j | \mathbf{x}_{\partial i})$$

and this can be used to identify $\mathbf{x}_{\partial i}$.

Reconstruction algorithm (Bresler, Mossel & Sly 2010) For each *i*, check $\begin{pmatrix} N \\ d \end{pmatrix}$ candidate neighborhoods ∂i . For each candidate ∂i , check condition on the remaining N - d - 1 nodes *j* Running time: $\sim N^{d+1}$ Inference in Computer Science and Systems Biology Part I — Other network reconstruction methods

The binary perceptron

■ The *perceptron* is an stylized model of a neuron and the simplest example of neural network (NN). The binary perceptron receives *x*₁,...,*x*_N (real valued) inputs and produces a binary output

$$\sigma = \operatorname{sign}\left(\sum_{i=1}^{N} w_i x_i\right) = \operatorname{sign}\left(\mathbf{w} \cdot \mathbf{x}\right)$$

- A perceptron is capable of *learning:* let's suppose we are given $\mathbf{x}^1, \ldots, \mathbf{x}^M$ patterns together with desired classification labels $\sigma^1, \ldots, \sigma^M$. The learning procedure consists in finding \mathbf{w} such that $\sigma^{\mu} = \text{sign}(\mathbf{w} \cdot \mathbf{x}^{\mu})$ for $\mu = 1, \ldots, M$
- This can be thought as the problem of finding the *separating plane*

The perceptron: generalizations and simplifications

- A slightly more general rule $\sigma = \text{sign} \left(\sum_{i=1}^{N} w_i x_i \theta \right)$ can be simply implemented as an extra dummy output $x_{N+1} = -1$
- We can assume $\sigma^{\tau} = +1$ for all τ ! Multiplying by σ^{τ} we get $1 = \sigma^{\tau} \sigma^{\tau} = \operatorname{sign} (\mathbf{w} \cdot (\sigma^{\tau} \mathbf{x}))$
- We will be interested in the following cases: $\mathbf{w} \in \mathbb{R}^N$ and $\mathbf{w} \in \{-1,1\}^N$ and $\mathbf{w} \in \{-q,\ldots,0,\ldots,q\}^N$
- \blacksquare We can assume that $\|\mathbf{x}^\tau\|=1$ since normalization doesn't affect classification

The online perceptron algorithm

Perceptron Algorithm

- **1** $w_0 = 0$
- **2** done = 0
- **B** while done = 0: **a** done = 1 **b** for $\tau = 1, ..., M$: **a** if $\mathbf{x}^{\tau} \cdot \mathbf{w}_t \le 0$ (mistake): $\mathbf{w}_{t+1} = \mathbf{w}_t + \mathbf{x}^{\tau}$ done = 0 $t \leftarrow t+1$

i.e. on any mistake, the algorithm greedily "helps" the classification of the missclassified pattern \mathbf{x}^{τ} , because $\mathbf{w}_{t+1} \cdot \mathbf{x}^{\tau} = (\mathbf{w}_t + \mathbf{x}^{\tau}) \cdot \mathbf{x}^{\tau} = \mathbf{w}_t \cdot \mathbf{x}^{\tau} + 1$

The Perceptron algorithm (analysis)

Assume there exists a classifier \mathbf{w}^* , i.e. $\mathbf{w}^* \cdot \mathbf{x}^{\tau} > 0$ for $\tau = 1, \dots, M$. Then the number of (mistake) events t must satisfy $t < \gamma^{-2}$

$$\gamma = \min_{\tau=1,...,M} \mathbf{x}^{\tau} \cdot \mathbf{w}^{*}$$

i.e. the algorithm must terminate in less than γ^{-2} iterations.

$$\begin{aligned} \mathbf{I} \quad \mathbf{w}_{t+1} \cdot \mathbf{w}^* &\geq \mathbf{w}_t \cdot \mathbf{w}^* + \gamma. \\ \text{Because } \mathbf{w}_{t+1} \cdot \mathbf{w}^* &= \mathbf{w}_t \cdot \mathbf{w}^* + \mathbf{x}^{\tau} \cdot \mathbf{w}^* \geq \mathbf{w}_t \cdot \mathbf{w}^* + \gamma \\ \mathbf{2} \quad \|\mathbf{w}_{t+1}\|^2 &\leq \|\mathbf{w}_t\|^2 + 1 \\ \text{Because } \|\mathbf{w}_{t+1}\|^2 &= \mathbf{w}_t \cdot \mathbf{w}_t + 2\mathbf{x}^{\tau} \cdot \mathbf{w}_t + \mathbf{x}^{\tau} \cdot \mathbf{x}^{\tau} \leq \|\mathbf{w}_t\|^2 + 1. \\ \text{This implies } \|\mathbf{w}_{t+1}\| &\leq \sqrt{t} \end{aligned}$$

Now after t mistakes, $t\gamma \leq \mathbf{w}_{t+1} \cdot \mathbf{w}^* \leq \|\mathbf{w}_{t+1}\| \leq \sqrt{t}$, thus $t \leq \gamma^{-2}$

I/O association as Bayesian Inference

We will just use the bayesian framework assuming that

- Data samples are formed by both input and output D = I, O
- The stochastic machine defines a stochastic rule P(O|S,I)
- S and I are independent

Then we can use Bayes:

$$P(S|I,O) = P(S,I,O)P(I,O)^{-1} = P(O|S,I)P(S)P(I)P(I,O)^{-1}$$

\$\approx P(O|S,I)P(S)\$

Similarly for $I^1, O^1, \ldots, I^M, O^M$ (assuming I^1, \ldots, I^M, S independent):

$$P\left(S|I^{1},O^{1},\ldots,I^{M},O^{M}\right) \propto \prod_{\mu=!}^{M} P\left(O^{\mu}|S,I^{\mu}\right) P\left(S\right)$$

Posterior distribution of binary perceptrons

Suppose we are given $I^1 = \mathbf{x}^1, O^1 = \sigma^1, \dots, I^M = \mathbf{x}^M, O^M = \sigma^M$ and we want to describe the posterior distribution for the binary perceptron $S = \mathbf{w}$

$$P(\mathbf{w}|\mathbf{x}^1, \sigma^1, \dots, \mathbf{x}^M, \sigma^M) \propto \prod_{\mu=1}^M P(\sigma^{\mu}|\mathbf{w}, \mathbf{x}^{\mu}) P(\mathbf{w})$$

The rule can be e.g. for $\sigma^{\mu} \in \{-1,1\}$ and $\mathbf{w}, \mathbf{x}^{\mu} \in \mathbb{R}^{N}$:

$$P(\sigma^{\mu}|\mathbf{w},\mathbf{x}^{\mu}) = \delta(\sigma^{\mu}; \operatorname{sign}(\mathbf{w}\cdot\mathbf{x}^{\mu}))$$

or more in general

$$P(\sigma^{\mu}|\mathbf{w},\mathbf{x}^{\mu}) = f(\sigma^{\mu};\mathbf{w}\cdot\mathbf{x}^{\mu})$$

Normally much easier to sample from I, O, given S than from a generic Boltzmann weight!

Posterior distribution as constraint satisfaction

- P(S) can be set to favour diluted classifiers S, e.g. $P(S) \propto \prod_i e^{\mu \delta(S_i,0)}$
- In fact, $P(S|I^1, O^1, \dots, I^M, O^M)$ can be thought as a *direct* model:

$$P(S|\mathbf{I},\mathbf{O}) \propto \prod_{\mu=1}^{M} \delta(O^{\mu}; \operatorname{sign}(S \cdot I^{\mu})) \prod_{i} e^{\mu \delta(S_{i},0)}$$

- And solved with mean-field approximations (e.g. Belief Propagation)
- Particularly simple if e.g. $S_i \in \{-q, \dots, 0, \dots, q\}$

Recurrent network

Suppose we have a binary network $\sigma_i \in \{-1, 1\}$, and $w_{ij} \in \{-q, \dots, 0, \dots, q\}$. Consider

$$P(\sigma|\mathbf{w}) \propto \prod_{i} \delta\left(\sigma_{i}; \operatorname{sign}\left(\sum_{j \neq i} w_{ji}\sigma_{j}\right)\right)$$

and dilution prior $P(\mathbf{w}) = \prod_{i \neq j} e^{\mu \delta(w_{ij}, 0)}$

$$P\left(\mathbf{w}|\sigma^{1},\ldots,\sigma^{\mu}\right) \propto \prod_{i} \left(\prod_{\mu=1}^{M} \delta\left(\sigma_{i}; \operatorname{sign}\left(\sum_{j\neq i} w_{ji}\sigma_{j}\right)\right) \prod_{j\neq i} e^{\mu\delta\left(w_{ij},0\right)}\right)$$

That is, the posterior distribution factorizes! N separate inference problems

For each *i*, BP can be used to find posterior statistics of the w_{ii} .

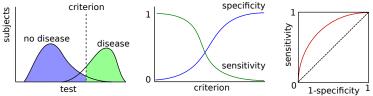
Insufficient data

- What to do if data is insufficient to infer a good model?
- ML/MAP are too risky: maybe the point of ML/MAP is not representative at all!
- In general we will be happier to get a small amount of sure information (e.g. a number of interactions that are present with high confidence) than a complete model with no poor confidence.
- How to measure performance (at least when we know the answer)? Something finer than correct/incorrect: ROC curves!

ROC curves

- ROC curves are thoroughly used in diagnostics
- Suppose we have a *test* which gives a scalar value $0 \le \alpha \le 1$ giving confidence of a certain disease.

Then depending on a given criterion value α_0 , we will predict *P* (disease) if $\alpha \ge \alpha_0$ and *N* (no disease) if $\alpha < \alpha_0$. How good is the test?



- Sensitivity=TP/(TP+FN)=TP/disease
- Specificity=TN/(TN+FP)=TN/no disease
- Area below the curve: discrimination. Probability for a random subject with disease to have α larger than that of a random subject without disease.

ROC curves for network inference

- Subject = edge
- With disease = present link, i.e. $e \in E$, $J_{ij} \neq 0$
- Without disease = absent link, i.e. $e \notin E$, $J_{ij} = 0$
- Criterion: e.g. M_{ij} , inferred $\left|J_{ij}^{ML}\right|$, inferred $\left|J_{ij}^{MAP}\right|$, $P\left(J_{ij} \neq 0 | \text{data}\right)$
- What criterion do we choose to have the best possible ROC curve?
- The best is to use $P(J_{ij} \neq 0 | \text{data})$ as criterion! Better expected ROC curve than MAP or ML estimate.

Things to read

- Yedidia, Weiss & Freeman, Belief propagation and its generalizations
 + variational interpretations
- David MacKay's book "Information Theory, Inference and Learning Algorithms"
- Jaynes paper on Maximum entropy
- Chow-Liu paper on inference on trees
- Mézard & Montanari's book
- Mezard & Mora's Susceptibility Propagation

The End

× ○ H: 15									8 00 0						
	8	A	R	辰	Ð	8		8	A	R	辰	Ð	5		
15	۵	R	R	盾		魚	R	商	R	()	A		儆	8	
A	Ð	颪		A	A	R	鹿	\mathbb{R}	5		12	A	R	Æ	
R		周	A	Æ	Æ	Е	R		\$	R	癒	Ν	Ε	8	
癥	R	æ	庖	Б	ß		A	S	B	Æ	N	5		15	0
R	۵	R	质		周	A	5	R	Æ	R		愈	R	Ε	0.08811 NS: 1.639e+50 Sn: 115
s	R		癒	B		Т	н	Е		臣	A	R	в	8	NS: 1
	5	R	R	盾	5	Ħ		A	12	Ð	8	Ε	B		.6394
6	Ð	Ř	Б	S		Е	Ν	D		6	质		重	8	+50 \$
A	Ð	颪		颪	R	8	ε	S	\$		凾	R	E	ß	n: 115
R		周	R	R	E	S	5		A	R	R	€	R	S	ία,
颪	R	Ð	卮	6	S		\$	ß	R	臣	Б	5		压	
E	۵	8	Æ		\$	ß	£	惪	Ε	8		颪	Ð	Ε	
s	R		5	ß	E	臣	Е	B		5	A	R	Ð	S	
	S	囼	A	Æ	Е	Ð		S	囼	A	Æ	Е	Ð		
Clean					Clear Fields										

