## MRF's, CRF's and Refining Localization

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## Program and Context

- CRF's and MRF's are important in semantic segmentation
- Work an interesting simple problem to set up
- Have a box on an object, but we'd like tighter boundaries
- What to do?
- Early (and very good) techniques
- Grab Cut
- Obj Cut
- Both use MRF/CRF models and inference
- cover that quickly


## Markov random field - formal

## Definition

Given an undirected graph $G=(V, E)$, a set of random variables $X=\left(X_{v}\right)_{v \in V}$ indexed by $V$ form a Markov random field with respect to $G$ if they satisfy the local Markov properties:

Pairwise Markov property: Any two non-adjacent variables are conditionally independent given all other variables:

$$
X_{u} \Perp X_{v} \mid X_{V \backslash\{u, v\}}
$$

Local Markov property: A variable is conditionally independent of all other variables given its neighbors:

$$
X_{v} \Perp X_{V \backslash \mathrm{~N}[v]} \mid X_{\mathrm{N}(v)}
$$

where $\mathrm{N}(v)$ is the set of neighbors of $v$, and $\mathrm{N}[v]=v \cup \mathrm{~N}(v)$ is the closed neighbourhood of $v$.

Global Markov property: Any two subsets of variables are conditionally independent given a separating subset:

$$
X_{A} \Perp X_{B} \mid X_{S}
$$

where every path from a node in $A$ to a node in $B$ passes through $S$.
The Global Markov property is stronger than the Local Markov property, which in turn is stronger than the Pairwise one.
${ }^{\text {[3] }}$ However, the above three Markov properties are equivalent for a positive probability. ${ }^{[4]}$

## MRF - First case for us



- The graph is a 2 D grid
- Each random variable is a binary random variable
- eg inside object, outside object
- In this case
$p(x) \propto \exp \left[\frac{1}{2} \sum_{i} \sum_{j} \operatorname{goodness}\left(x_{i}, x_{j}\right)\right]$

Look at Ch15 of AML for some examples, BUT that uses different inference procedures and has 1,-1 labels. I'm using Greig; Porteous; Seheult notation (see web page for paper)

## Notice

- If the goodness of a pair is high, p is higher
- Because these are binary, we can simplify
- We want:
- better for neighbors to agree than disagree
- the goodness for both 0 is the same as for both 1
- Can then simplify

$$
p(x) \propto \exp \left[\frac{1}{2} \sum_{i} \sum_{j} \operatorname{goodness}\left(x_{i}, x_{j}\right)\right]
$$

- To get
$p(x) \propto \exp \left[\frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \beta_{i j}\left\{x_{i} x_{j}+\left(1-x_{i}\right)\left(1-x_{j}\right)\right\}\right]$


## Important

- We want:
- better for neighbors to agree than disagree
- the goodness for both 0 is the same as for both 1
- This means



## First model

- At each pixel, there is an unknown binary label - $0=$ out, $1=$ in
- These binary labels form an MRF
- where it is cheaper to agree than to disagree
- At each pixel, there are measurements
- conditioned on the label
- details to follow
- Q: how do we get the MAP set of labels?


## Model

- At each pixel we have observations y
- yields likelihood

$$
l(y \mid x)=\prod_{i=1}^{n} f\left(y_{i} \mid x_{i}\right)=\prod_{i=1}^{n} f\left(y_{i} \mid 1\right)^{x_{i}} f\left(y_{i} \mid 0\right)^{1-x_{i}}
$$

- what is f ? (later)
- write $\quad \lambda_{i}=\ln \left\{f\left(y_{i} \mid 1\right) / f\left(y_{i} \mid 0\right)\right\}$
- Then

$$
\begin{aligned}
\log p(x \mid y)= & \sum_{i=1}^{n} \lambda_{i} x_{i}+\frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \beta_{i j}\left\{x_{i} x_{j}+\left(1-x_{i}\right)\left(1-x_{j}\right)\right\} \\
& +K
\end{aligned}
$$

## To obtain MAP estimate

- Maximise

$$
\sum_{i=1}^{n} \lambda_{i} x_{i}+\frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \beta_{i j}\left\{x_{i} x_{j}+\left(1-x_{i}\right)\left(1-x_{j}\right)\right\}
$$

- But how?
- blank search won't do it (why?)
- In this special case, graph cut works


## Graph cut (quick but clean)

Consider a capacitated network comprising $n+2$ vertices, being a source $s$, a sink $t$ and the $n$ pixels. There is a directed edge ( $s, i$ ) from $s$ to pixel $i$ with capacity $c_{s i}=\lambda_{i}$, if $\lambda_{i}>0$; otherwise, there is a directed edge $(i, t)$ from $i$ to $t$ with capacity $c_{i t}=-\lambda_{i}$. There is an undirected edge (i,j) between two internal vertices (pixels) $i$ and $j$ with capacity $c_{i j}=\beta_{i j}$ if the corresponding pixels are neighbours.


## Graph cut (quick but clean)

For any binary image $x=\left(x_{1}, \ldots, x_{n}\right)$ let $B=\{s\} \cup\left\{i: x_{i}=1\right\}$ and $W=$ $\left\{i: x_{i}=0\right\} \cup\{t\}$ define a two-set partition of the network vertices and put

$$
C(x)=\sum_{k \in B} \sum_{l \in W} c_{k l} .
$$



## Graph cut (quick but clean)

The set of edges with a vertex in $B$ and a vertex in $W$ is called a cut and $C(x)$ is called the capacity of the cut.

It ${ }^{t}$ is readily seen that $C(x)$ may be written

$$
C(x)=\sum_{i=1}^{n} x_{i} \max \left(0,-\lambda_{i}\right)+\sum_{i=1}^{n}\left(1-x_{i}\right) \max \left(0, \lambda_{i}\right)+\frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \beta_{i j}\left(x_{i}-x_{j}\right)^{2}
$$

which differs from $-L(x \mid y)$ by a term which does not depend on $x$;


## Graph cut, II

- SO
- set up the graph as described, and do a min-cut
- this is polynomial
- Ifs, ands, buts
- this only works in the case it is cheaper to agree than to disagree
- more general case, it's max cut which isn't funny at all
- this only works for the binary case
- but approximations for some multilabel cases are very good
- More details
- there are *many* min-cut algorithms with different complexities
- adapted to different types of problem
- significant literature on best min-cut algorithm for vision applications
- we'll ignore - search github


## Grab Cut

- Originally for matting
- extracting an object from an image
- Process
- user places box
- grabcut segments intended object
- user perhaps iterates with strokes, etc.
- For us:

- segments using graph cuts
- clever iterative model of interior/exterior
- extremely simple shape prior on object



## Simplest case: grey level image

Their paper [Boykov and Jolly 2001] addresses the segmentation of a monochrome image, given an initial trimap $T$. The image is an array $\mathbf{z}=\left(z_{1}, \ldots, z_{n}, \ldots, z_{N}\right)$ of grey values, indexed by the (single) index $n$. The segmentation of the image is expressed as an array of "opacity" values $\underline{\alpha}=\left(\alpha_{1}, \ldots, \alpha_{N}\right)$ at each pixel. Generally $0 \leq \alpha_{n} \leq 1$, but for hard segmentation $\alpha_{n} \in\{0,1\}$, with 0 for background and 1 for foreground. The parameters $\underline{\theta}$ describe image foreground and background grey-level distributions, and consist of histograms of grey values:

$$
\begin{equation*}
\underline{\theta}=\{h(z ; \alpha), \alpha=0,1\}, \tag{1}
\end{equation*}
$$

one for background and one for foreground. The histograms are assembled directly from labelled pixels from the respective trimap regions $T_{B}, T_{F}$. (Histograms are normalised to sum to 1 over the grey-level range: $\int_{z} h(z ; \alpha)=1$.)

## Grey level image, II

An energy function $\mathbf{E}$ is defined so that its minimum should correspond to a good segmentation, in the sense that it is guided both by the observed foreground and background grey-level histograms and that the opacity is "coherent", reflecting a tendency to solidity of objects. This is captured by a "Gibbs" energy of the form:

$$
\begin{equation*}
\mathbf{E}(\underline{\alpha}, \underline{\theta}, \mathbf{z})=U(\underline{\alpha}, \underline{\theta}, \mathbf{z})+V(\underline{\alpha}, \mathbf{z}) . \tag{2}
\end{equation*}
$$

The data term $U$ evaluates the fit of the opacity distribution $\underline{\alpha}$ to the data $\mathbf{z}$, given the histogram model $\underline{\theta}$, and is defined to be:

$$
\begin{equation*}
U(\underline{\alpha}, \underline{\theta}, \mathbf{z})=\sum_{n}-\log h\left(z_{n} ; \alpha_{n}\right) \tag{3}
\end{equation*}
$$

The smoothness term can be written as

$$
\begin{equation*}
V(\underline{\alpha}, \mathbf{z})=\gamma \sum_{(m, n) \in \mathbf{C}} \operatorname{dis}(m, n)^{-1}\left[\alpha_{n} \neq \alpha_{m}\right] \exp -\beta\left(z_{m}-z_{n}\right)^{2}, \tag{4}
\end{equation*}
$$

where $[\phi]$ denotes the indicator function taking values 0,1 for a predicate $\phi, \mathbf{C}$ is the set of pairs of neighboring pixels, and where $\operatorname{dis}(\cdot)$ is the Euclidean distance of neighbouring pixels. This energy

## Notice

$$
\begin{aligned}
& \sum_{i=1}^{n} \lambda_{i} x_{i}+\frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \beta_{i j}\left\{x_{i} x_{j}+\left(1-x_{i}\right)\left(1-x_{j}\right)\right\} \\
& V(\underline{\alpha}, \mathbf{z})=\gamma \sum_{(m, n) \in \mathbf{C}} \operatorname{dis}(m, n)^{-1}\left[\alpha_{n} \neq \alpha_{m}\right] \exp -\beta\left(z_{m}-z_{n}\right)^{2}, \\
& U(\underline{\alpha}, \underline{\theta}, \mathbf{z})=\sum_{n}-\log h\left(z_{n} ; \alpha_{n}\right) .
\end{aligned}
$$

They're minimizing, and GPS are maximizing; this means they use a cost (not goodness) for disagreeing (not agreeing)

## Improving this

- Where does trimap come from?
- start with
- inside: a bunch of pixels in "deep interior" of box
- outside: a bunch of pixels outside box
- Histograms for color images are clumsy
- too big
- Initial trimap is messy
- reestimate using segmentation


## Replace histograms

- Use mixture of normals
- have some interior, some exterior pixels
- build mixture of normal model for each case
- AML ch 9 if you've forgotten
- now you can compute $\mathrm{p}(\mathrm{yl} 1)$, etc. from this


## Re-estimation

- Use initial trimap to make GMM
- Segment with graph cut
- Now you have a trimap
- Re-estimate GMMs, and iterate


## An alternative strategy

- Variational inference
- High level:
- come up with simpler model that is "most like" intractable model
- extract information from that
- Currently:
- chose x_i (each 0 or 1 ) to maximize expression below

$$
\begin{aligned}
\log p(x \mid y)= & \sum_{i=1}^{n} \lambda_{i} x_{i}+\frac{1}{2} \sum_{i=1}^{n} \sum_{j=1}^{n} \beta_{i j}\left\{x_{i} x_{j}+\left(1-x_{i}\right)\left(1-x_{j}\right)\right\} \\
& +K
\end{aligned}
$$

## New setup

- H - hidden variables, 1 or -1
- used to write x
- X - observations

$$
\log p(H \mid X)=\sum_{i}^{\text {known }} \stackrel{\downarrow}{w}_{i} H_{i}+\sum_{i j}^{q_{i j}} H_{i}^{\text {known }} H_{j}-\log Z
$$

- notice the $-1,1$ trick


### 15.2 VARIATIONAL INFERENCE

We could just ignore intractable models, and stick to tractable models. This isn't a good idea, because intractable models are often quite natural. The discrete Markov random field model of an image is a fairly natural model. Image labels should depend on pixel values, and on neighboring labels. It is better to try and deal with the intractable model. One really successful strategy for doing so is to choose a tractable parametric family of probability models $Q(H ; \theta)$, then adjust $\theta$ to find parameter values $\hat{\theta}$ that represent a distribution that is "close" in the right sense to $P(H \mid X)$. One then extracts information from $Q(H ; \hat{\theta})$. This process is known as variational inference. What is remarkable is that (a) it is possible to find a $Q(H ; \hat{\theta})$ without too much fuss and (b) information extracted from this distribution is often accurate and useful.
15.2.1 The KL Divergence

Assume we have two probability distributions $P(X)$ and $Q(X)$. A measure of their similarity is the KL-divergence (or sometimes Kullback-Leibler divergence) written

$$
\mathbb{D}(P \| Q)=\int P(X) \log \frac{P(X)}{Q(X)} d X
$$

(you've clearly got to be careful about zeros in $P$ and $Q$ here). This likely strikes you as an odd measure of similarity, because it isn't symmetric. It is not the case that $\mathbb{D}(P \| Q)$ is the same as $\mathbb{D}(Q \| P)$, which means you have to watch your P 's and Q's. Furthermore, some work will demonstrate that it does not satisfy the triangle inequality, so KL divergence lacks two of the three important properties of a metric.

KL divergence has some nice properties, however. First, we have

$$
\mathbb{D}(P \| Q) \geq 0
$$

with equality only if $P$ and $Q$ are equal almost everywhere (i.e. except on a set of measure zero).

> Remember this: The KL divergence measures the similarity of two probability distributions. It is always non-negative, and is only zero if the two distributions are the same. However, it is not symmetric.

## KL divergence and Maximum likelihood

Second, there is a suggestive relationship between KL divergence and maximum likelihood. Assume that $X_{i}$ are IID samples from some unknown $P(X)$, and we wish to fit a parametric model $Q(X \mid \theta)$ to these samples. This is the usual situation we deal with when we fit a model. Now write $H(P)$ for the entropy of $P(X)$, defined by

$$
H(P)=-\int P(X) \log P(X) d x=-\mathbb{E}_{P}[\log P]
$$

The distribution $P$ is unknown, and so is its entropy, but it is a constant. Now we can write

$$
\mathbb{D}(P \| Q)=\mathbb{E}_{P}[\log P]-\mathbb{E}_{P}[\log Q]
$$

## KL divergence and Maximum Likelihood

Then

$$
\begin{aligned}
\mathcal{L}(\theta)=\sum_{i} \log Q\left(X_{i} \mid \theta\right) \approx \int P(X) \log Q(X \mid \theta) d X & =\mathbb{E}_{P(X)}[\log Q(X \mid \theta)] \\
& =-H(P)-\mathbb{D}(P \| Q)(\theta)
\end{aligned}
$$

Equivalently, we can write

$$
\mathcal{L}(\theta)+\mathbb{D}(P \| Q)(\theta)=-H(P)
$$

Recall $P$ doesn't change (though it's unknown), so $H(P)$ is also constant (though unknown). This means that when $\mathcal{L}(\theta)$ goes up, $\mathbb{D}(P \| Q)(\theta)$ must go down. When $\mathcal{L}(\theta)$ is at a maximum, $\mathbb{D}(P \| Q)(\theta)$ must be at a minimum. All this means that, when you choose $\theta$ to maximize the likelihood of some dataset given $\theta$ for a parametric family of models, you are choosing the model in that family with smallest KL divergence from the (unknown) $P(X)$.

### 15.2.2 The Variational Free Energy

We have a $P(H \mid X)$ that is hard to work with (usually because we can't evaluate $P(X))$ and we want to obtain a $Q(H)$ that is "close to" $P(H \mid X)$. A good choice of "close to" is to require that

$$
\mathbb{D}(Q(H) \| P(H \mid X))
$$

is small. Expand the expression for KL divergence, to get

$$
\begin{aligned}
\mathbb{D}(Q(H) \| P(H \mid X)) & =\mathbb{E}_{Q}[\log Q]-\mathbb{E}_{Q}[\log P(H \mid X)] \\
& =\mathbb{E}_{Q}[\log Q]-\mathbb{E}_{Q}[\log P(H, X)]+\mathbb{E}_{Q}[\log P(X)] \\
& =\mathbb{E}_{Q}[\log Q]-\mathbb{E}_{Q}[\log P(H, X)]+\log P(X)
\end{aligned}
$$

which at first glance may look unpromising, because we can't evaluate $P(X)$. But $\log P(X)$ is fixed (although unknown). Now rearrange to get

$$
\begin{aligned}
\log P(X) & =\mathbb{D}(Q(H) \| P(H \mid X))-\left(\mathbb{E}_{Q}[\log Q]-\mathbb{E}_{Q}[\log P(H, X)]\right) \\
& =\mathbb{D}(Q(H) \| P(H \mid X))-\mathrm{E}_{Q} .
\end{aligned}
$$

Here

$$
\mathrm{E}_{Q}=\left(\mathbb{E}_{Q}[\log Q]-\mathbb{E}_{Q}[\log P(H, X)]\right)
$$

is referred to as the variational free energy. We can't evaluate $\mathbb{D}(Q(H) \| P(H \mid X))$. But, because $\log P(X)$ is fixed, when $\mathrm{E}_{Q}$ goes down, $\mathbb{D}(Q(H) \| P(H \mid X))$ must go down too. Furthermore, a minimum of $\mathrm{E}_{Q}$ will correspond to a minimum of $\mathbb{D}(Q(H) \| P(H \mid X))$. And we can evaluate $\mathrm{E}_{Q}$.

## Variational Inference

We now have a strategy for building approximate $Q(H)$. We choose a family of approximating distributions. From that family, we obtain the $Q(H)$ that minimises $\mathrm{E}_{Q}$ (which will take some work). The result is the $Q(H)$ in the family that minimizes $\mathbb{D}(Q(H) \| P(H \mid X))$. We use that $Q(H)$ as our approximation to $P(H \mid X)$, and extract whatever information we want from $Q(H)$.

- Questions:
- what $\mathrm{Q}(\mathrm{H})$ ?
- hard, case by case basis; essentially, so that calculations go through
- How to minimize?
- straightforward (long, dull) calculation
- (AML Ch15 for easiest example)


## Variational Inference

We want to construct a $Q(H)$ that approximates the posterior for a Boltzmann machine. We will choose $Q(H)$ to have one factor for each hidden variable, so $Q(H)=q_{1}\left(H_{1}\right) q_{2}\left(H_{2}\right) \ldots q_{N}\left(H_{N}\right)$. We will then assume that all but one of the terms in $Q$ are known, and adjust the remaining term. We will sweep through the terms doing this until nothing changes.

The $i^{\prime}$ th factor in $Q$ is a probability distribution over the two possible values of $H_{i}$, which are 1 and -1 . There is only one possible choice of distribution. Each $q_{i}$ has one parameter $\pi_{i}=P\left(\left\{H_{i}=1\right\}\right)$. We have

$$
q_{i}\left(H_{i}\right)=\left(\pi_{i}\right)^{\frac{\left(1+H_{i}\right)}{2}}\left(1-\pi_{i}\right)^{\frac{\left(1-H_{i}\right)}{2}} .
$$

Notice the trick; the power each term is raised to is either 1 or 0 , and I have used this trick as a switch to turn on or off each term, depending on whether $H_{i}$ is 1 or -1 . So $q_{i}(1)=\pi_{i}$ and $q_{i}(-1)=\left(1-\pi_{i}\right)$. This is a standard, and quite useful, trick. We wish to minimize the variational free energy, which is

$$
\mathrm{E}_{Q}=\left(\mathbb{E}_{Q}[\log Q]-\mathbb{E}_{Q}[\log P(H, X)]\right)
$$

