

- PhD student since 2008
- Main focus is on construction of discrete Markov random field models from training images (Toftaker and Tjelmeland, 2012).
- In the fall of 2010 I was in Aalborg, working with professor Jesper Møller on anisotropic spatial point patterns (Møller and Toftaker, 2012).
- Now we work on another approach to construct discrete models from training images.
- The thesis will be finished in November 2012.

# Construction of Markov random field prior models.

Håkon Toftaker

NTNU

February 8, 2012

- Construct realistic Markov random field prior models for rock types in petroleum reservoirs.
- Models should be constructed from training data such as these.



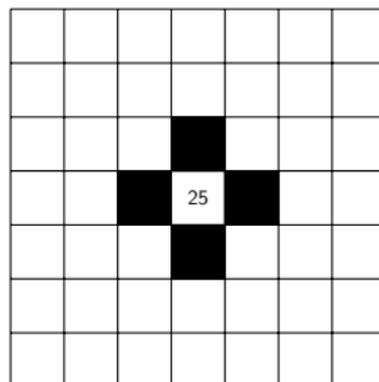
# Markov random fields

- The Markov random field  $x = (x_1, \dots, x_n)$  is a set of binary variables defined on a lattice of  $n$  nodes

$x_1$	$x_2$	$x_3$	$x_4$	$x_5$	$x_6$	$x_7$
$x_8$	$x_9$	$x_{10}$	$x_{11}$	$x_{12}$	$x_{13}$	$x_{14}$
$x_{15}$	$x_{16}$	$x_{17}$	$x_{18}$	$x_{19}$	$x_{20}$	$x_{21}$
$x_{22}$	$x_{23}$	$x_{24}$	$x_{25}$	$x_{26}$	$x_{27}$	$x_{28}$
$x_{29}$	$x_{30}$	$x_{31}$	$x_{32}$	$x_{33}$	$x_{34}$	$x_{35}$
$x_{36}$	$x_{37}$	$x_{38}$	$x_{39}$	$x_{40}$	$x_{41}$	$x_{42}$
$x_{43}$	$x_{44}$	$x_{45}$	$x_{46}$	$x_{47}$	$x_{48}$	$x_{49}$

# Markov random fields

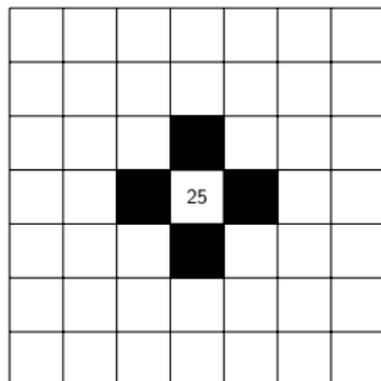
- The Markov random field  $x = (x_1, \dots, x_n)$  is a set of binary variables defined on a lattice of  $n$  nodes
- To each node  $k$  we associate a neighborhood  $\partial k$



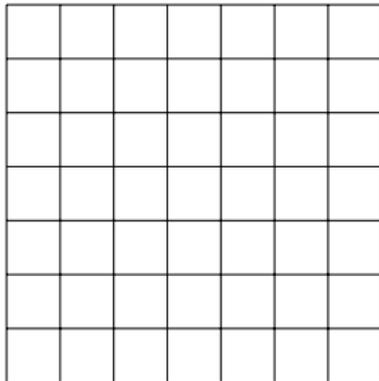
# Markov random fields

- The Markov random field  $x = (x_1, \dots, x_n)$  is a set of binary variables defined on a lattice of  $n$  nodes
- To each node  $k$  we associate a neighborhood  $\partial k$
- The probability distribution of  $x$  has a Markov property, which means

$$p_\theta(x_k | x_{-k}) = p_\theta(x_k | x_{\partial k})$$



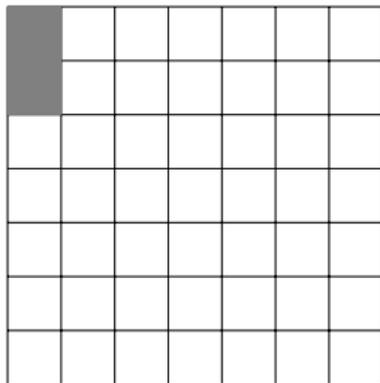
- A clique is defined as a set of nodes, which are all neighbors of each other.



- A clique is defined as a set of nodes, which are all neighbors of each other.
- The corresponding joint distribution is

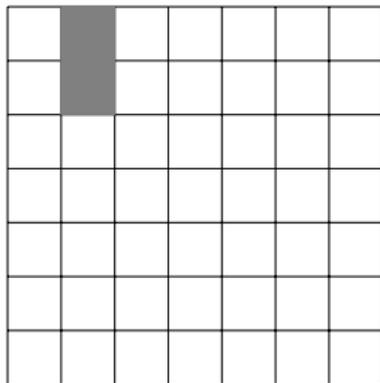
$$p_{\theta}(x) = c(\theta) \exp \left\{ - \sum_{C \in \mathcal{C}} V(x_C, \theta) \right\}$$

$$= c(\theta) \exp \{ -U_{\theta}(x) \}.$$



- A clique is defined as a set of nodes, which are all neighbors of each other.
- The corresponding joint distribution is

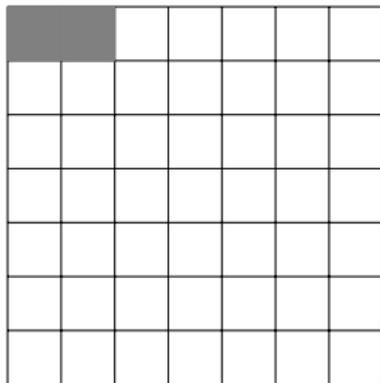
$$p_{\theta}(x) = c(\theta) \exp \left\{ - \sum_{C \in \mathcal{C}} V(x_C, \theta) \right\}$$
$$= c(\theta) \exp \{ -U_{\theta}(x) \}.$$



- A clique is defined as a set of nodes, which are all neighbors of each other.
- The corresponding joint distribution is

$$p_{\theta}(x) = c(\theta) \exp \left\{ - \sum_{C \in \mathcal{C}} V(x_C, \theta) \right\}$$

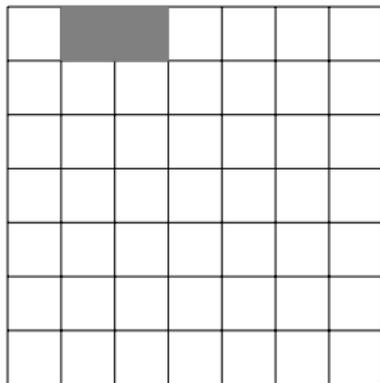
$$= c(\theta) \exp \{ -U_{\theta}(x) \}.$$



- A clique is defined as a set of nodes, which are all neighbors of each other.
- The corresponding joint distribution is

$$p_{\theta}(x) = c(\theta) \exp \left\{ - \sum_{C \in \mathcal{C}} V(x_C, \theta) \right\}$$

$$= c(\theta) \exp \{ -U_{\theta}(x) \}.$$



- Here  $c(\theta)$  is a normalizing constant

$$c(\theta) = \sum_x \exp \{ -U_{\theta}(x) \},$$

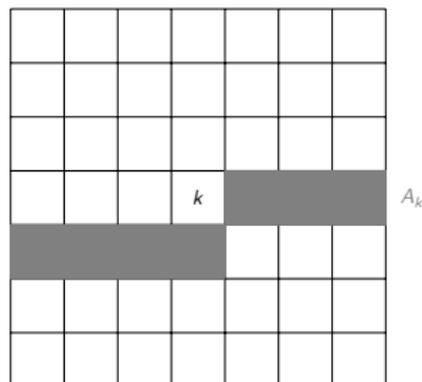
and  $U_{\theta}(x)$  is called an energy function.

# Forward-backward algorithm

- By a forward-backward algorithm (Friel and Rue, 2005) we can sum out the components of  $x$ , and sequentially obtain

$$p_{\theta}(x_k | x_{k+1:n}) = p_{\theta}(x_k | x_{A_k})$$

and  $c(\theta)$ .



# Forward-backward algorithm

- By a forward-backward algorithm (Friel and Rue, 2005) we can sum out the components of  $x$ , and sequentially obtain

$$p_{\theta}(x_k | x_{k+1:n}) = p_{\theta}(x_k | x_{A_k})$$

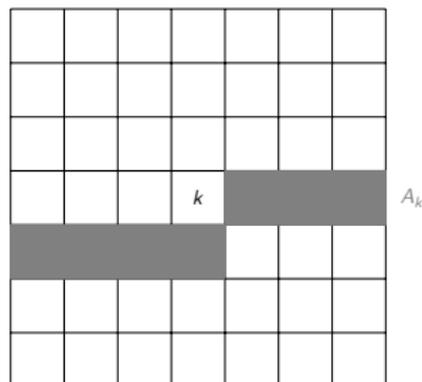
and  $c(\theta)$ .

- The probability distribution is available as

$$p_{\theta}(x) = p_{\theta}(x_n) \prod_{k=1}^{n-1} p_{\theta}(x_k | x_{A_k}),$$

and it is simple to sample from  $p_{\theta}(x)$ .

- Computational complexity is  $2^{|A_k|+1} = 2^{7+1}$



# The approximation

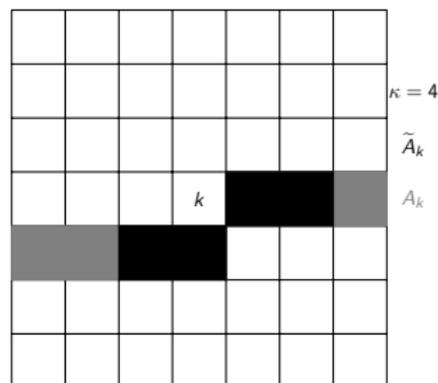
- Austad and Tjelmeland, 2011, develop a computationally feasible approximation of the above algorithm.

# The approximation

- Austad and Tjelmeland, 2011, develop a computationally feasible approximation of the above algorithm.
- An approximate factorization is obtained

$$\tilde{p}_\theta(x) = \tilde{p}_\theta(x_n) \prod_{k=1}^{n-1} \tilde{p}_\theta(x_k | x_{\tilde{A}_k}),$$

where  $\tilde{A}_k \subseteq A_k$ , and  $|\tilde{A}_k| < \kappa$ .



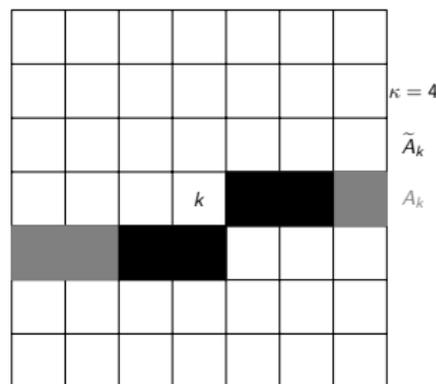
# The approximation

- Austad and Tjelmeland, 2011, develop a computationally feasible approximation of the above algorithm.
- An approximate factorization is obtained

$$\tilde{p}_\theta(x) = \tilde{p}_\theta(x_n) \prod_{k=1}^{n-1} \tilde{p}_\theta(x_k | x_{\tilde{A}_k}),$$

where  $\tilde{A}_k \subseteq A_k$ , and  $|\tilde{A}_k| < \kappa$ .

- The computational complexity is reduced to  $2^{\kappa+1}$



# Multi-grid MRF

- A large neighborhood is required to capture the structure in the training images.
- A large neighbourhood increases the size of  $A_k$  and increases the error of the approximation for a given value of  $\kappa$ .
- We split the lattice into  $T$  sublattices  $S_1, \dots, S_T$ .

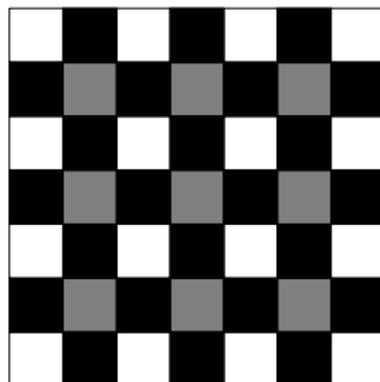


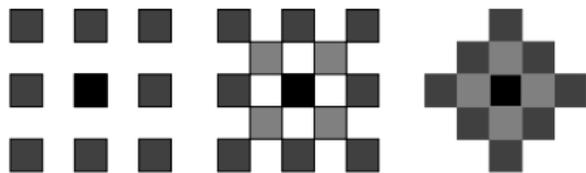
Figure:  $T = 3$

- The joint distribution for  $x$  is specified by the marginal of  $x_{S_1}$  and the conditionals of  $x_{S_t}$  given  $x_{S_{1:t-1}}$

$$\begin{aligned}
 p_{\theta}(x) &= p_{\theta_1}(x_{S_1}) \prod_{t=2}^T p_{\theta_t}(x_{S_t} | x_{S_{1:t-1}}) \\
 &= c(\theta_1) \exp\{-U_{\theta_1}(x_{S_1})\} \\
 &\quad \prod_{t=2}^T c(\theta_t, x_{S_{1:t-1}}) \exp\{-U_{\theta_t}(x_{S_1}, x_{S_{1:t-1}})\}
 \end{aligned}$$

- The energy function has the form

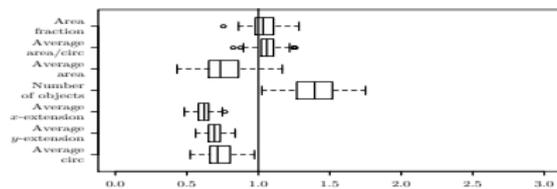
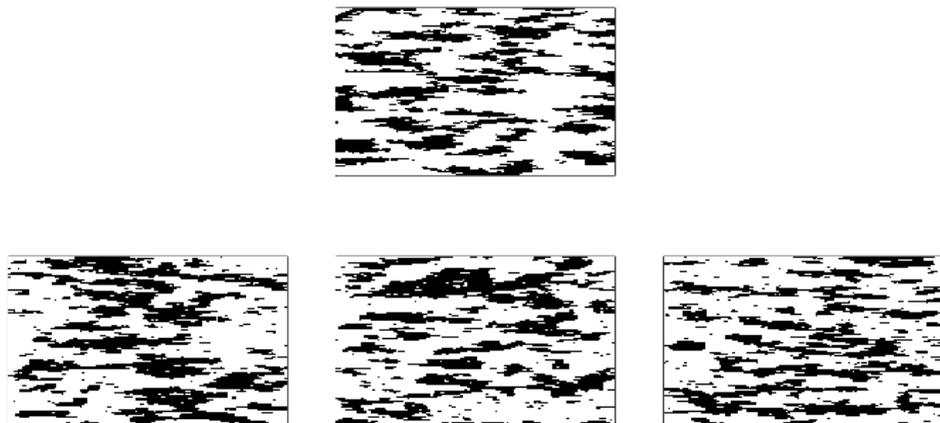
$$U_{\theta_t}(x_{S_t}, x_{S_{1:t-1}}) = U_{\theta_{t,1:9}}^1(x_{S_t}) + \sum_{i \in S_t} U_{\theta_{t,10:19}}^2(x_i, x_{B_i})$$

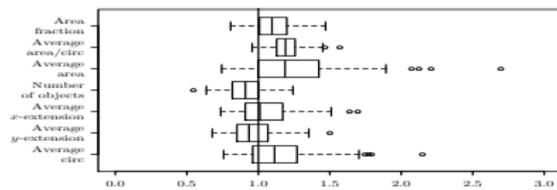
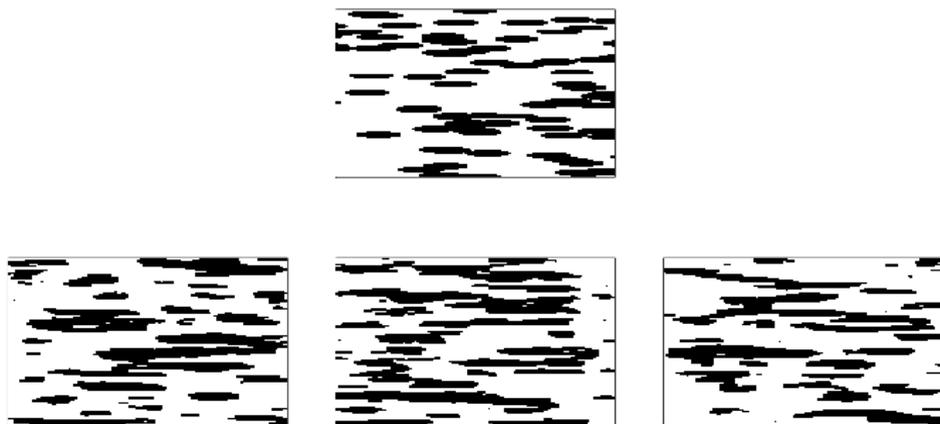


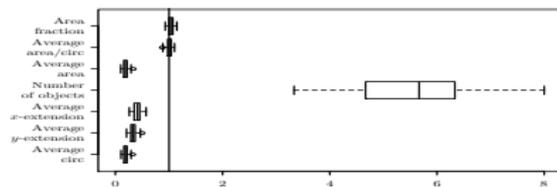
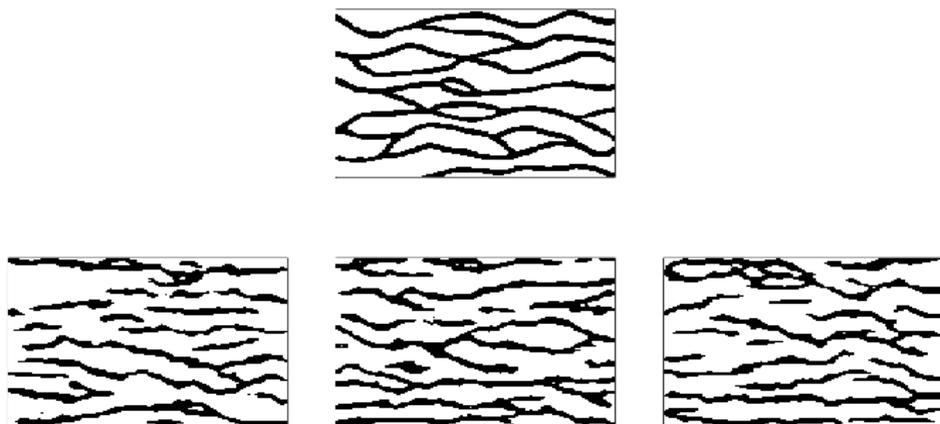
**Figure:** The dark gray nodes in  $S_t$  which are neighbors of the black node. The light gray are nodes from  $S_{1:t-1}$  in  $B_i$

- Approximate simulation from  $p_\theta(x)$  is done sequentially by first drawing  $x_{S_1}$  from  $\tilde{p}_{\theta_1}(x_{S_1})$  and then  $x_{S_2}$  from  $\tilde{p}_{\theta_2}(x_{S_2}|x_{S_1})$  and so on.
- In the examples we have used  $T = 7$  sublattices.

# Unconditional simulation







# Conditional simulation

- Conditional simulation is complicated by the unknown normalizing constants  $c(\theta_t, x_{S_{1:t-1}})$ .
- To solve this problem we apply the approximate forward-backward algorithm to  $p_{\theta_t}(x_{S_t} | x_{S_{1:t-1}})$ , and obtain  $\tilde{c}(\theta_t, x_{S_{1:t-1}})$
- The resulting approximation is

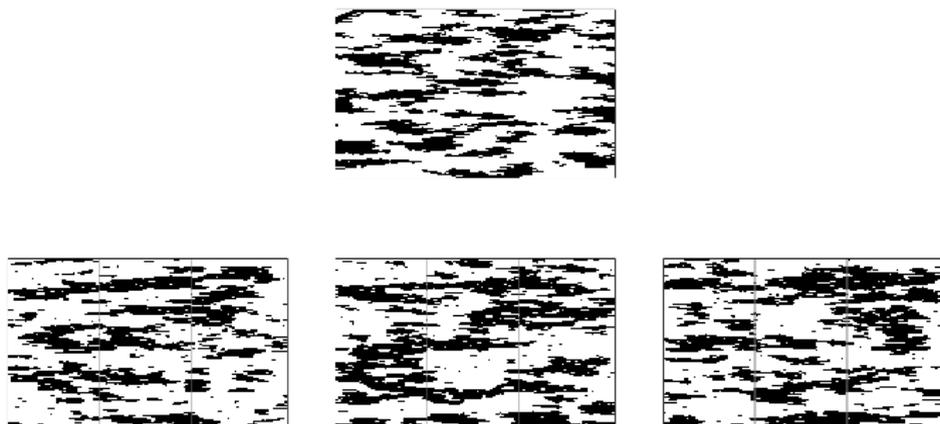
$$\tilde{p}_{\theta}(x) = \tilde{c}(\theta_1) \exp\{-U_{\theta_1}(x_{S_1})\} \prod_{t=2}^T \tilde{c}(\theta_t, x_{S_{1:t-1}}) \exp\{-U_{\theta_t}(x_{S_{1:t}})\},$$

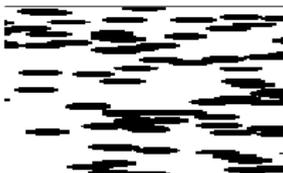
- let  $z$  denote a vector of observed quantities which is related to  $x$  by a likelihood function  $h(z|x)$ .
- For the approximation  $\tilde{p}_\theta(x)$  we get a corresponding approximations to  $p_\theta(x|z)$

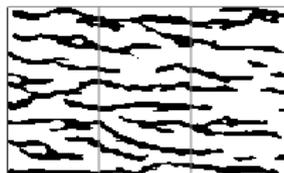
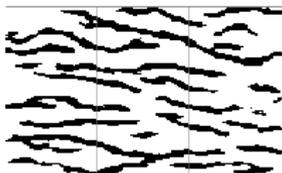
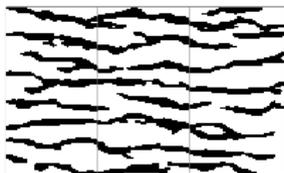
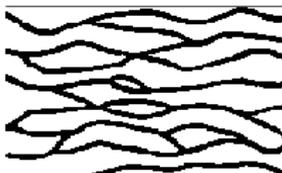
$$\tilde{p}_\theta(x|z) \propto \tilde{p}_\theta(x)h(z|x)$$

- The approximate distribution  $\tilde{p}_\theta(x|z)$  is an MRF, and we can apply the approximate forward-backward algorithm again.

# Conditioning on 2 wells







- The method seems to be quite flexible.
- Conditional simulation is straight forward if the likelihood is easy to compute.
- A Generalization to 3D is simple, but would probably be computationally unfeasible.
- More than 2 colors is also straight forward but computational complexity increases fast.