# Markov Random Fields in Image Segmentation

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#### Overview

- Segmentation as pixel labeling
- Probabilistic approach
  - Markov Random Field (MRF)
  - □ Gibbs distribution & Energy function
- Energy minimization
  - Simulated Annealing
  - Markov Chain Monte Carlo (MCMC) sampling
- Example MRF model & Demo
- Parameter estimation (EM)
- More complex models

## Segmentation as a Pixel Labelling Task

- 1. Extract features from the input image
  - □ Each pixel s in the image has a feature vector
  - For the whole image, we have

$$f = \{\vec{f}_s : s \in S\}$$

- 2. Define the set of labels  $\Lambda$ 
  - $\square$  Each pixel s is assigned a label  $\omega_s \in \Lambda$
  - □ For the whole image, we have

$$\omega = \{\omega_s, s \in S\}$$

- For an  $N \times M$  image, there are  $|A|^{NM}$  possible labelings.
  - Which one is the right segmentation?







#### Probabilistic Approach, MAP

- Define a <u>probability measure</u> on the set of all possible labelings and select the most likely one.
- $P(\omega \mid f)$  measures the probability of a labelling, given the observed feature f
- Our goal is to find an optimal labeling  $\hat{\omega}$  which maximizes  $P(\omega \mid f)$
- This is called the <u>Maximum a Posteriori</u> (MAP) estimate:

$$\hat{\omega}^{MAP} = \arg\max_{\omega \in \Omega} P(\omega \mid f)$$

likelihood

prior



#### **Bayesian Framework**

By Bayes Theorem, we have

$$P(\omega \mid f) = \frac{P(f \mid \omega)P(\omega)}{P(f)} \propto P(f \mid \omega)P(\omega)$$

- $\blacksquare P(f)$  is constant
- We need to define  $P(\omega)$  and  $P(f \mid \omega)$  in our model

#### Why MRF Modelization?

- In real images, regions are often homogenous; neighboring pixels usually have similar properties (intensity, color, texture, ...)
- Markov Random Field (MRF) is a probabilistic model which captures such contextual constraints
- Well studied, strong theoretical background
- Allows MCMC sampling of the (hidden) underlying structure → Simulated Annealing



#### What is MRF?

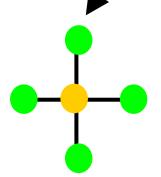
- To give a formal definition for Markov Random Fields, we need some basic building blocks
  - Observation Field and (hidden) Labeling Field
  - □ Pixels and their Neighbors
  - Cliques and Clique Potentials
  - □ Energy function
  - □ Gibbs Distribution

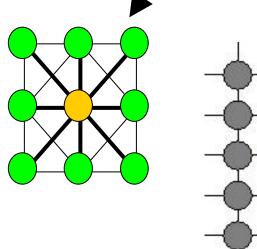


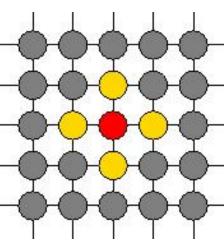
#### **Definition – Neighbors**

For each pixel, we can define some surrounding pixels as its neighbors.

Example : 1<sup>st</sup> order neighbors and 2<sup>nd</sup> order neighbors







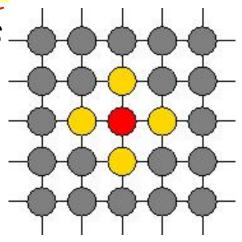


#### **Definition – MRF**

- The labeling field X can be modeled as a Markov Random Field (MRF) if
  - 1. For all  $\omega \in \Omega : P(X = \omega) > 0$
  - 2. For every  $s \in S$  and  $\omega \in \Omega$ :

$$P(\omega_s \mid \omega_r, r \neq s) = P(\omega_s \mid \omega_r, r \in N_s)$$

 $N_s$  denotes the neighbors of pixel s





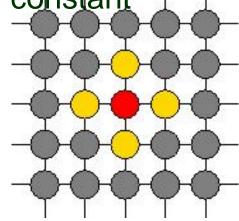
#### **Hammersley-Clifford Theorem**

■ The Hammersley-Clifford Theorem states that a random field is a MRF if and only if  $P(\omega)$  follows a Gibbs distribution.

$$P(\omega) = \frac{1}{Z} \exp(-U(\omega)) = \frac{1}{Z} \exp(-\sum_{c \in C} V_c(\omega))$$

• where  $Z = \sum_{\omega \in \Omega} \exp(-U(\omega))$  is a normalization constant

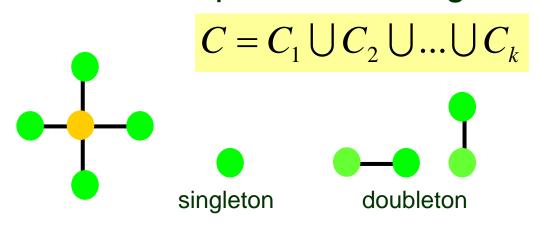
This theorem provides us an easy way of defining MRF models via <u>clique potentials</u>.

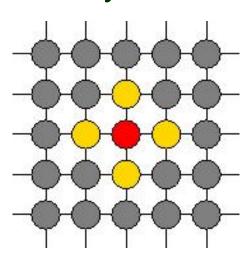




#### **Definition – Clique**

- A subset  $C \subseteq S$  is called a <u>clique</u> if every pair of pixels in this subset are neighbors.
- A clique containing n pixels is called n<sup>th</sup> order clique, denoted by  $C_n$ .
- The set of cliques in an image is denoted by



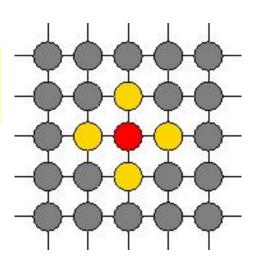




#### **Definition – Clique Potential**

- For each clique c in the image, we can assign a value  $V_c(\omega)$  which is called <u>clique potential</u> of c, where  $\omega$  is the configuration of the labeling field
- The sum of potentials of all cliques gives us the energy  $U(\omega)$  of the configuration  $\omega$

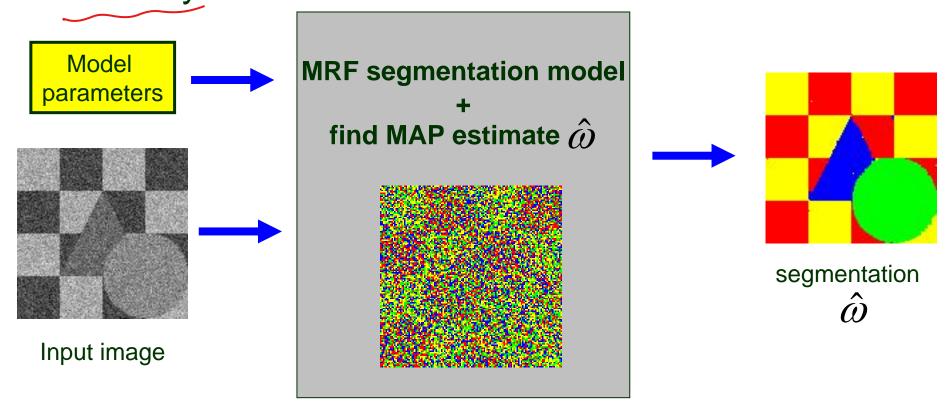
$$U(\omega) = \sum_{c \in C} V_c(\omega) = \sum_{i \in C_1} V_{C_1}(\omega_i) + \sum_{(i,j) \in C_2} V_{C_2}(\omega_i, \omega_j) + \dots$$





# Segmentation of grayscale images: A simple MRF model

Construct a segmentation model where regions are formed by spatial clusters of pixels with similar intensity:





# MRF segmentation model

Pixel labels (or classes) are represented by Gaussian distributions:

$$P(f_s \mid \omega_s) = \frac{1}{\sqrt{2\pi\sigma_{\omega_s}}} \exp\left(-\frac{(f_s - \mu_{\omega_s})^2}{2\sigma_{\omega_s}^2}\right)$$

- Clique potentials:
  - □ **Singleton**: proportional to the likelihood of features given  $\omega$ :  $log(P(f \mid \omega))$ .
  - Doubleton: favours similar labels at neighbouring pixels smoothness prior

$$V_{c_2}(i,j) = \beta \delta(\omega_i, \omega_j) = \begin{cases} -\beta & \text{if} & \omega_i = \omega_j \\ +\beta & \text{if} & \omega_i \neq \omega_j \end{cases}$$

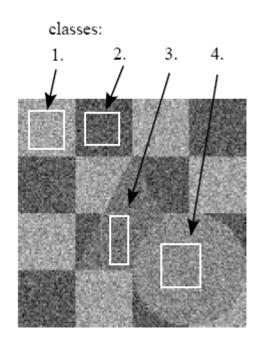
Cliques

As  $\beta$  increases, regions become more homogenous



# Model parameters

- Doubleton potential β
  - □ less dependent on the input →
    - can be fixed a priori
- Number of labels (|\Lambda|)
  - □ Problem dependent →
    - usually given by the user or
    - inferred from some higher level knowledge
- Each label  $\lambda \in \Lambda$  is represented by a Gaussian distribution  $N(\mu_{\lambda}, \sigma_{\lambda})$ :
  - estimated from the input image



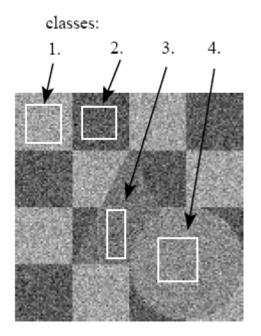


# Model parameters

The class statistics (mean and variance) can be estimated via the *empirical mean* and variance:

$$\forall \lambda \in \Lambda : \qquad \mu_{\lambda} = \frac{1}{|S_{\lambda}|} \sum_{s \in S_{\lambda}} f_{s},$$
$$\sigma_{\lambda}^{2} = \frac{1}{|S_{\lambda}|} \sum_{s \in S_{\lambda}} (f_{s} - \mu_{\lambda})^{2}$$

- $\square$  where  $S_{\lambda}$  denotes the set of pixels in the training set of class  $\lambda$
- a training set consists in a representative region selected by the user





# **Energy function**

Now we can define the energy function of our MRF model:

our MRF model: 
$$U(\omega) = \sum_{s} \left( \log(\sqrt{2\pi}\sigma_{\omega_{s}}) + \frac{(f_{s} - \mu_{\omega_{s}})^{2}}{2\sigma_{\omega_{s}}^{2}} \right) + \sum_{s,r} \beta \delta(\omega_{s}, \omega_{r})$$

■ Recall:

$$P(\omega \mid f) = \frac{1}{Z} \exp(-U(\omega)) = \frac{1}{Z} \exp(-\sum_{c \in C} V_c(\omega))$$

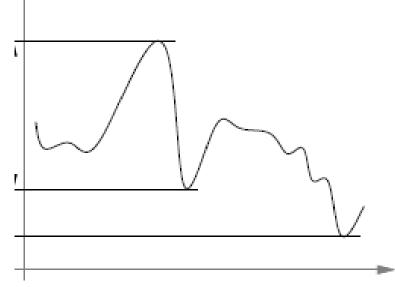
Hence

$$\hat{\omega}^{MAP} = \arg \max_{\omega \in \Omega} P(\omega \mid f) = \arg \min_{\omega \in \Omega} U(\omega)$$



# Optimization

- Problem reduced to the minimization of a non-convex energy function
  - Many local minima
- Gradient descent?
  - □ Works only if we have a good initial segmentation
- Simulated Annealing
  - □ Always works (at least in theory)



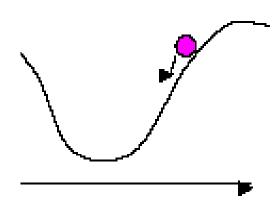


# ICM (~Gradient descent) [Besag86]

- ① Start at a "good" initial configuration  $\omega^0$  and set k=0.
- ② For each configuration which differs at most in one element from the current configuration ω<sup>k</sup> (they are denoted by N<sub>ωk</sub>), compute the energy U(η) (η ∈ N<sub>ωk</sub>).
- ③ From the configurations in N<sub>ωk</sub>, select the one which has a minimal energy:

$$\omega^{k+1} = \arg\min_{\eta \in \mathcal{N}_{\omega^k}} U(\eta). \tag{6}$$

④ Goto Step ② with k = k + 1 until convergence is obtained (for example, the energy change is less than a certain threshold).





# Simulated Annealing

- ① Set k = 0 and initialize  $\omega$  randomly. Choose a sufficiently high initial temperature  $T = T_0$ .
- ② Construct a trial perturbation  $\eta$  from the current configuration  $\omega$  such that  $\eta$  differs only in one element from  $\omega$ .
- ③ (Metropolis criteria) Compute  $\Delta U = U(\eta) U(\omega)$  and accept  $\eta$  if  $\Delta U < 0$  else accept with probability  $\exp(-\Delta U/T)$  (analogy with thermodynamics):

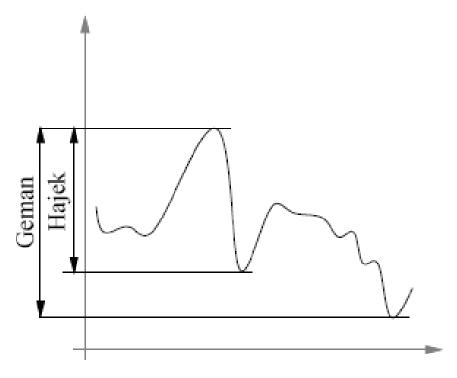
$$\omega = \begin{cases} \eta & \text{if } \Delta U \leq 0, \\ \eta & \text{if } \Delta U > 0 \text{ and } \xi < \exp(-\Delta U/T), \\ \omega & \text{otherwise} \end{cases}$$
 (4)

where  $\xi$  is a uniform random number in [0,1).

4 Decrease the temperature:  $T = T_{k+1}$  and goto Step 2 with k = k+1 until the system is frozen.



# Temperature Schedule



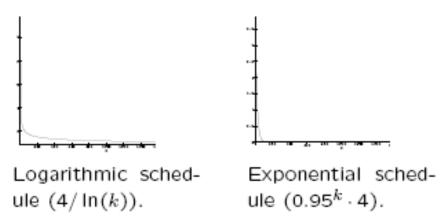
$$T_k \ge \frac{\Gamma}{\ln(k)}$$
 (8)

with

$$\Gamma > \max_{\omega \in \Omega} U(\omega) - \min_{\omega \in \Omega} U(\omega) \tag{9}$$



# Temperature Schedule



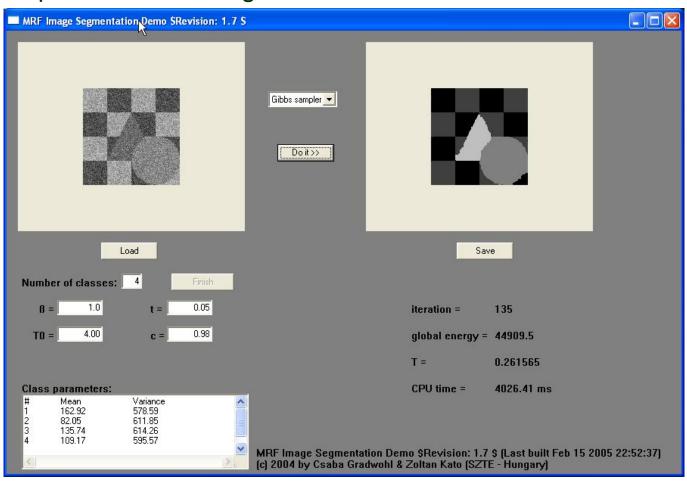
- Initial temperature: set it to a relatively low value (~4)→ faster execution
  - □ must be high enough to allow random jumps at the beginning!
- Schedule:  $T_{k+1} = c \cdot T_k, \quad k = 0, 1, 2, ...$
- Stopping criteria:
  - ☐ Fixed number of iterations
  - □ Energy change is less than a threshols



# Demo

Download from:

http://www.inf.u-szeged.hu/~kato/software/



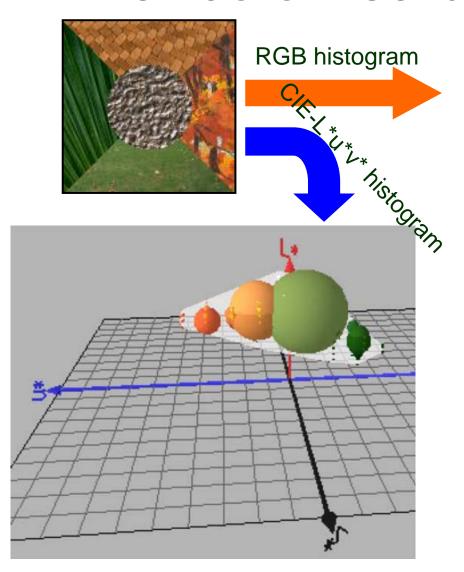


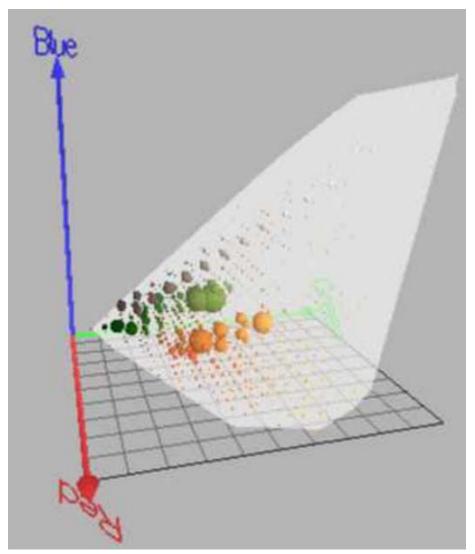
# Summary

- Design your model carefully
  - Optimization is just a tool, do not expect a good segmentation from a wrong model
- What about other than graylevel features
  - Extension to color is relatively straightforward



# What color features?



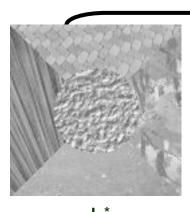


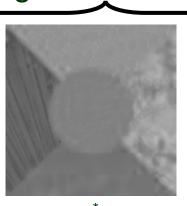


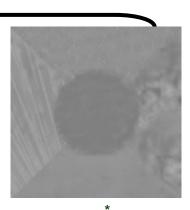
#### **Extract Color Feature**

- We adopt the CIE-L\*u\*v\* color space because it is perceptually uniform.
  - Color difference can be measured by Euclidean distance of two color vectors.
- We convert each pixel from RGB space to CIE-L\*u\*v\* space →
  - □ We have 3 color feature images











# Color MRF segmentation model

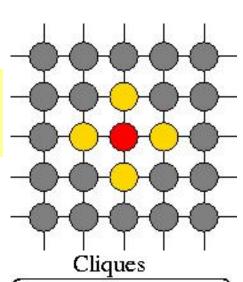
Pixel labels (or classes) are represented by three-variate Gaussian distributions:

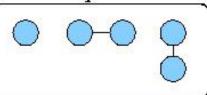
$$P(f_{s} \mid \omega_{s}) = \frac{1}{\sqrt{(2\pi)^{n} \mid \Sigma_{\omega_{s}} \mid}} \exp(-\frac{1}{2}(\vec{f}_{s} - \vec{u}_{\omega_{s}})\Sigma_{\omega_{s}}^{-1}(\vec{f}_{s} - \vec{u}_{\omega_{s}})^{T})$$

- Clique potentials:
  - □ **Singleton**: proportional to the likelihood of features given  $\omega$ :  $log(P(f \mid \omega))$ .
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$$V_{c_2}(i,j) = \beta \delta(\omega_i, \omega_j) = \begin{cases} -\beta & \text{if} & \omega_i = \omega_j \\ +\beta & \text{if} & \omega_i \neq \omega_j \end{cases}$$

As  $\beta$  increases, regions become more homogenous







# Summary

- Design your model carefully
  - Optimization is just a tool, do not expect a good segmentation from a wrong model
- What about other than graylevel features
  - Extension to color is relatively straightforward
- Can we segment images without user interaction?
  - Yes, but you need to estimate model parameters automatically (EM algorithm)



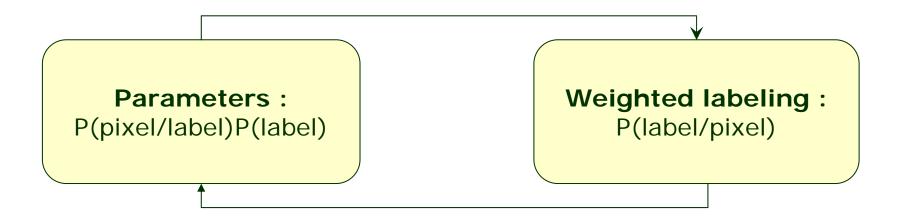
# Incomplete data problem

- Supervised parameter estimation
  - we are given a labelled data set to learn from
    - e.g. somebody manually assigned labels to pixels
- How to proceed without labelled data?
  - Learning from incomplete data
  - □ Standard solution is an iterative procedure called *Expectation-Maximization* 
    - Assigns labels and estimates parameters simultaneously
    - Chicken-Egg problem



# EM principles: The two steps

**E Step**: For each pixel, use parameters to compute probability distribution



**M Step**: Update the estimates of parameters based on weighted (or "soft") labeling



## The basic idea of EM

- Each of the E and M steps is straightforward assuming the other is solved
  - Knowing the label of each pixel, we can estimate the parameters
    - Similar to supervised learning (hard vs. soft labeling)
  - Knowing the parameters of the distributions, we can assign a label to each pixel
    - by Maximum Likelihood i.e. using the singleton energies only without pairwise interactions



# Parameter estimation via EM

- Basically, we will fit a mixture of Gaussian to the image histogram
  - □ We know the number of labels  $|\Lambda|$  = number of mixture components
- At each pixel, the complete data includes
  - □ The observed feature f<sub>s</sub>
  - $\square$  Hidden pixel labels  $I_s$  (a vector of size  $|\Lambda|$ )
    - specifies the contribution of the pixel feature to each of the labels – i.e. a soft labeling



## Parameter estimation via EM

■ E step: recompute I<sub>s</sub> at each pixel s:

$$\mathbf{l_s^i} = P(\lambda \mid \mathbf{f_s}) = \frac{P(\mathbf{f_s} \mid \lambda)P(\lambda)}{\sum_{\lambda \in \Lambda} P(\mathbf{f_s} \mid \lambda)P(\lambda)}$$

■ M step: update Gaussian parameters for

each label  $\lambda$ :

$$P(\lambda) = \frac{\sum_{s \in S} P(\lambda \mid \mathbf{f_s})}{\mid S \mid}, \quad \mu_{\lambda} = \frac{\sum_{s \in S} P(\lambda \mid \mathbf{f_s}) \mathbf{f_s}}{\sum_{s \in S} P(\lambda \mid \mathbf{f_s})}, \dots$$



# Summary

- Design your model carefully
  - Optimization is just a tool, do not expect a good segmentation from a wrong model
- What about other than graylevel features
  - □ Extension to color is relatively
- Can we segment images without user interaction?
  - Yes, but you need to estimate model parameters automatically (EM algorithm)
- What if we do not know  $|\Lambda|$ ?
  - □ Fully automatic segmentation requires
    - Modeling of the parameters AND
    - a more sophisticated sampling algorithm (Reversible jump MCMC)



# MRF+RJMCMC vs. JSEG





RJMCMC (17 min)

#### JSEG (Y. Deng, B.S.Manjunath: PAMI'01):

- 1. **color quantization**: colors are quantized to several representing classes that can be used to differentiate regions in the image.
- **2. spatial segmentation**: A region growing method is then used to segment the image.





# Benchmark results using the Berkeley Segmentation Dataset







JSEG RJMCMC



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  - Yes, but you need to estimate model parameters automatically (EM algorithm)
- What if we do not know  $|\Lambda|$ ?
  - □ Fully automatic segmentation requires
    - Modeling of the parameters AND
    - a more sophisticated sampling algorithm (Reversible jump MCMC)
- Can we segment more complex images?
  - □ Yes but you need a more complex MRF model



### Objectives

- Combine different segmentation cues:
  - □ Color & Texture [ICPR2002,ICIP2003]
  - □ Color & Motion [ACCV2006,ICIP2007]
  - □...?
- How humans do it?
  - □ Multiple cues are perceived simultaneously and then they are integrated by the human visual system [Kersten *et al. An. Rev. Psych.* 2004]
  - □ Therefore different image features has to be handled in a parallel fashion.
- We attempt to develop such a model in a Markovian framework

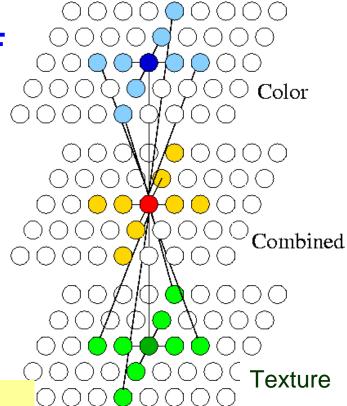


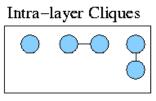
### Multi-Layer MRF Model: Neighborhood & Interactions

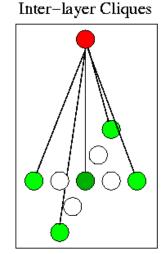
- $\bullet$  is modeled as a MRF
  - □ Layered structure
  - □ "Soft" interaction between features
- $\rightarrow$   $P(\omega \mid f)$  follows a Gibbs distribution
  - Clique potentials define the local interaction strength
- MAP  $\Leftrightarrow$  Energy minimization  $(U(\omega))$

Hammersley - Clifford Theorem:

$$P(\omega) = \frac{\exp(-U(\omega))}{Z} = \frac{\exp(-\sum_{C} V_{C}(\omega))}{Z}$$







**Model** ⇔ **Definition of clique potentials** 

### Texture Layer: MRF model

- We extract two type of texture features
  - ☐ <u>Gabor feature</u> is good at discriminating strongordered textures
  - MRSAR feature is good at discriminating weakordered (or random) textures
  - □ The number of texture feature images depends on the size of the image and other parameters.
    - Most of these doesn't contain useful information →
  - Select feature images with high discriminating power.
- MRF model is similar to the color layer model.



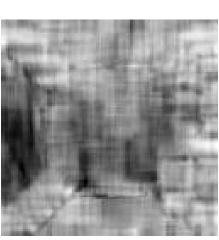
# **Examples of Texture Features**

**Gabor features:** 

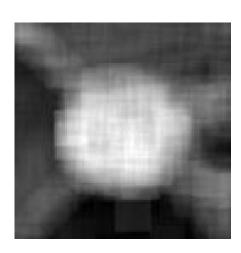


MRSAR features:







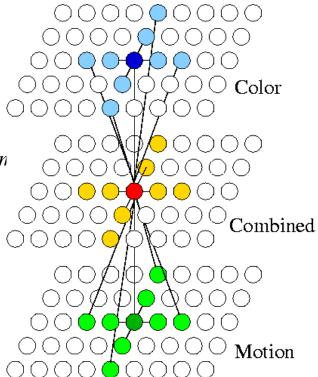


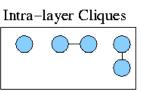


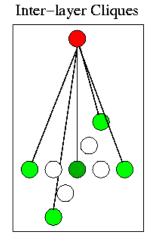


# Combined Layer: Labels

- A label on the combined layer consists of a pair of color and texture/motion labels such that  $\eta_s = \langle \eta_s^c, \eta_s^m \rangle$  where  $\eta_s^c \in \Lambda^c$  and  $\eta_s^m \in \Lambda^n$
- The number of possible classes is  $L^c \times L^m$
- The combined layer selects the most likely ones.









#### Combined Layer: Singleton potential

Controls the number of classes:

$$V_s(\eta_s) = R((10N_{\eta_s})^{-3} + P(L))$$

- $\square N_{\eta_s}$  is the percentage of labels belonging to class  $\eta_s$
- $\square$  L is the number of classes present on the combined layer.
- Unlikely classes have a few pixels → they will be penalized and removed to get a lower energy
- $\blacksquare$  P(L) is a log-Gaussian term:
  - □ Mean value is a guess about the number of classes,
  - Variance is the confidence.



#### Combined Layer: **Doubleton** potential

- Preferences are set in this order:
  - 1. Similar color and motion/texture labels
  - Different color and motion/texture labels
  - Similar color (resp. motion/texture) and different motion/texture (resp. color) labels
    - These are contours visible only at one feature layer.

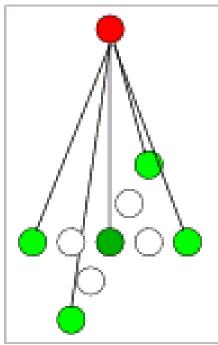
$$\mathcal{S}(\eta_s, \eta_r) = \begin{cases} -\alpha & \text{if } \eta_s^c = \eta_r^c, \eta_s^m = \eta_r^m \\ 0 & \text{if } \eta_s^c \neq \eta_r^c, \eta_s^m \neq \eta_r^m \\ +\alpha & \text{if } \eta_s^c \neq \eta_r^c, \eta_s^m = \eta_r^m \\ & \text{or } \eta_s^c = \eta_r^c, \eta_s^m \neq \eta_r^m \end{cases}$$



## Inter-layer clique potential

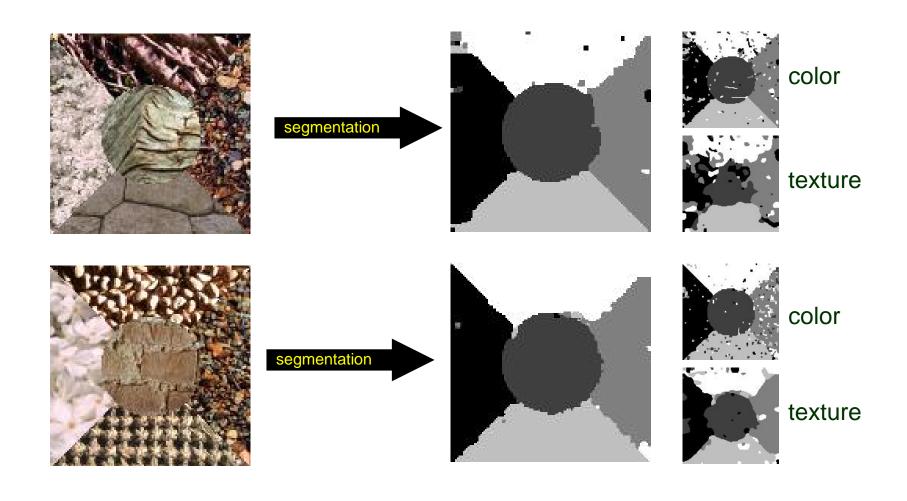
- Five pair-wise interactions between a feature and combined layer
- Potential is proportional to the difference of the singleton potentials at the corresponding feature layer.
  - $\square$  Prefers  $\omega_s$  and  $\eta_s$  having the same label, since they represent the labeling of the same pixel
  - $\square$  Prefers  $\omega_s$  and  $\eta_r$  having the same label, since we expect the combined and feature layers to be homogenous





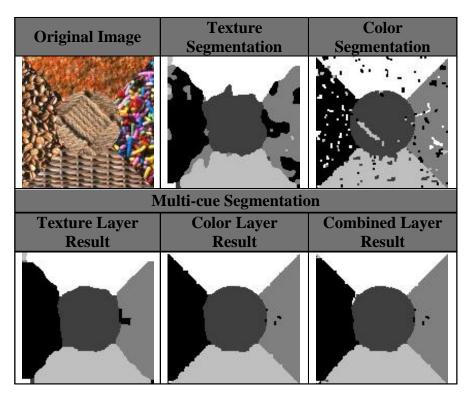


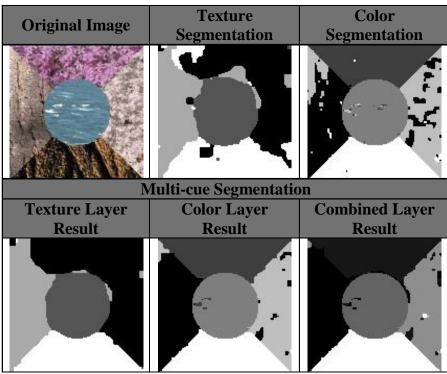
# Color Textured Segmentation





# Color Textured Segmentation



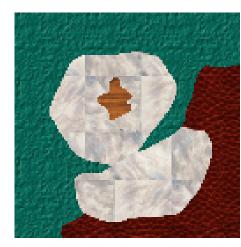


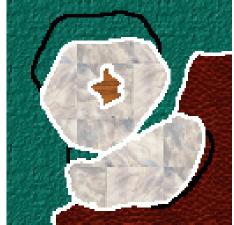


# Color & Motion Segmentation















#### References

Visit

http://www.inf.u-szeged.hu/~kato/