
Markov Random Fields in Image Segmentation

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Zoltan Kato

*Image Processing and Computer Graphics Dept.
University of Szeged
Szeged 6720
Hungary
kato@inf.u-szeged.hu*

Josiane Zerubia

*INRIA Sophia Antipolis-Mediterranee
Sophia Antipolis
06902 Cedex
France
Josiane.Zerubia@inria.fr*

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Markov Random Fields in Image Segmentation

Zoltan Kato¹ and Josiane Zerubia²

¹ *Image Processing and Computer Graphics Dept., University of Szeged,
Arpad ter 2, Szeged, 6720, Hungary, kato@inf.u-szeged.hu*

² *INRIA Sophia Antipolis-Mediterranee, 2004 Route des Lucioles, Sophia
Antipolis, 06902 Cedex, France, Josiane.Zerubia@inria.fr*

Abstract

This monograph gives an introduction to the fundamentals of Markovian modeling in image segmentation as well as a brief overview of recent advances in the field. Segmentation is considered in a common framework, called image labeling, where the problem is reduced to assigning labels to pixels. In a probabilistic approach, label dependencies are modeled by Markov random fields (MRF) and an optimal labeling is determined by Bayesian estimation, in particular maximum a posteriori (MAP) estimation. The main advantage of MRF models is that prior information can be imposed locally through clique potentials. The primary goal is to demonstrate the basic steps to construct an easily applicable MRF segmentation model and further develop its multiscale and hierarchical implementations as well as their combination in a multilayer model. MRF models usually yield a non-convex energy function. The minimization of this function is crucial in order to find the most likely segmentation according to the MRF model. Besides classical optimization algorithms, like simulated annealing or

deterministic relaxation, we also present recently introduced graph cut-based algorithms. We briefly discuss the possible parallelization techniques of simulated annealing, which allows efficient implementation on, e.g., GPU hardware without compromising convergence properties of the algorithms. While the main focus of this monograph is on generic model construction and related energy minimization methods, many sample applications are also presented to demonstrate the applicability of these models in real life problems such as remote sensing, biomedical imaging, change detection, and color- and motion-based segmentation. In real-life applications, parameter estimation is an important issue when implementing completely data-driven algorithms. Therefore some basic procedures, such as expectation-maximization, are also presented in the context of color image segmentation.

Note: A sample implementation of the most important segmentation algorithms is available in grey scale at http://dx.doi.org/10.1561/20000000035_demogray and in color at http://dx.doi.org/10.1561/20000000035_democolor.

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Dedication

“To the memory of my mother” Zoltan Kato

“To the memory of my beloved sister Elise who passed away in August
2012” Josiane Zerubia

1

Introduction

An image processing system involves a sensing device (usually a camera) and computer algorithms to interpret the picture. The term *image* (more precisely, *monochrome image*) refers to a two-dimensional light intensity function whose value at any point is proportional to the brightness (*gray-level*) of the image at that point [70]. A *digital image* is a discretized image both in spatial coordinates and in brightness. It is usually represented as a two-dimensional matrix, the elements of such a digital array are called pixels. The digitized image is the starting point of any kind of computer analysis. In some applications, the sensing device may be more specific responding to other forms of light: infrared imaging, photon emission tomography, radar imaging [182], ultrasonic imaging, etc.

Many image processing tasks deal directly with raw pixel data involving image compression [2], restoration [35, 64, 91, 219, 220, 223], edge detection [65, 200, 219, 220, 223], segmentation [51, 52, 60, 61, 74, 83, 98, 115, 195, 196, 221], texture analysis [43, 66, 122], motion detection [90, 213], optical flow and motion analysis [87, 90, 167], etc. Most of these problems can be formulated in a general framework, called *image labeling*, where we associate a label to each pixel from a finite set. The meaning of this label depends on the problem that we

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are trying to solve. For image restoration, it means a gray-level; for edge detection, it means the presence or the direction of an edge; for image segmentation, it means a region; etc. The problem here is how to choose a label for a pixel, which is *optimal* in a certain sense. Herein, we deal with a statistical approach of *labeling*. In real scenes, neighboring pixels usually have similar features (intensity, color, texture, etc). In a probabilistic framework, such regularities are well expressed mathematically by Markov random fields. In this survey, we will focus on the fundamental problem of image segmentation using Markovian models.

1.1 Image Segmentation

The primary goal of any segmentation algorithm is to divide the domain R of the input image into the disjoint parts R_i such that they belong to distinct objects in the scene. The solution of this problem sometimes requires high level knowledge about the shape and appearance of the objects under investigation [46, 123, 183, 202]. In many applications, however, such information is not available or impractical to use. Hence low-level features of the surface patches are used for the segmentation process [9, 141, 224]. Herein, we are interested in the latter approach. In either case, we have to summarize all relevant information in a model which is then adjusted to fit the image data.

One broadly used class of models is the so called *cartoon model*, which has been extensively studied from both probabilistic [64] and variational [19, 163, 169] viewpoints. The model assumes that the real world scene consists of a set of regions whose observed low-level features change slowly, but across the boundary between them, these features change abruptly. What we want to infer is a *cartoon* ω consisting of a simplified, abstract version of the input image \mathcal{I} : regions R_i have a constant value (called a *label* in our context) and the discontinuities between them form a curve Γ — the contour. The pair (ω, Γ) specifies a *segmentation*. Region based methods are mainly focused on ω while edge based methods try to determine Γ directly.

Taking the probabilistic approach, one usually wants to come up with a *probability measure* on the set Ω of all possible segmentations of \mathcal{I} and then select the one with the highest probability. Note that

Ω is finite, although huge. A widely accepted standard, also motivated by the human visual system [121, 162], is to construct this probability measure in a Bayesian framework [37, 161, 214]: We shall assume that we have a set of observed (Y) and hidden (X) random variables. In our context, any observed value $y \in Y$ represents the low-level features used for partitioning the image, and the hidden entity $x \in X$ represents the segmentation itself. First, we have to quantify how well any occurrence of x fits y . This is expressed by the probability distribution $P(y|x)$ — the *imaging model*. Second, we define a set of properties that any segmentation x must possess regardless the image data. These are described by $P(x)$, the *prior*, which tells us how well any occurrence x satisfies these properties. Factoring these distributions and applying the Bayes theorem gives us the *posterior* distribution $P(x|y) \propto P(y|x)P(x)$. Note that the constant factor $1/P(y)$ has been dropped as we are only interested in \hat{x} which *maximizes* the posterior, that is, the maximum a posteriori (MAP) estimate of the hidden field X .

The models of the above distributions also depend on certain parameters that we denote by Θ . Supervised segmentation assumes that these parameters are either known or a set of joint realizations of the hidden field X and observations Y (called a *training set*) is available [64, 205]. This is known in statistics as the *complete data* problem which is generally easier to solve than the *incomplete case* [37]. Although the prior knowledge of the parameters is a strong assumption, supervised methods are still useful alternatives when working in a controlled environment. Many industrial applications, like quality inspection of agricultural products [166], fall into this category. In the unsupervised case, however, we know neither Θ nor X . This is called the *incomplete data* problem where both Θ and X have to be inferred from the only observable entity Y . Hence our MAP estimation problem becomes $(\hat{x}, \hat{\Theta}) = \arg \max_{x, \Theta} P(x, \Theta|y)$. *Expectation Maximization* (EM) [48] and its variants (Stochastic EM [33, 149], Gibbsian EM [36]), as well as *Iterated Conditional Expectation* (ICE) [30, 108] are widely used to solve such problems. It is important to note, however, that these methods calculate a local maximum [37].

Due to the difficulty of estimating the number of pixel classes (or clusters), unsupervised algorithms often suppose that this parameter

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is *known a priori* [68, 77, 141, 145, 149]. When the number of pixel classes is also being estimated, the unsupervised segmentation problem may be treated as a *model selection* problem over a combined model space [102, 202, 203].

1.2 Markov Random Fields

In the early 20th century, mostly inspired by the Ising model [170], a new type of stochastic process appeared in the theory of probability, called *Markov random field* (MRF). MRFs rapidly became a broadly used tool in a variety of problems, not only in statistical mechanics. The use of MRFs in image processing became popular with the seminal paper of S. Geman and D. Geman [64] in 1984, but its first use in the domain dates to the early 70s [16, 215]. Here, we give a brief introduction to the theory of MRFs [39, 54, 57, 79, 125, 144, 160, 184, 214].

1.2.1 The Ising Model

Following Ising [10, 69, 170], we consider a sequence, $0, 1, 2, \dots, n$ on the line. At each point, there is a small spin which is either *up* or *down* at any given moment (see Figure 1.1). Now, we define a probability measure on the set Ω of all possible configurations $\omega = (\omega_0, \omega_1, \dots, \omega_n)$. In this context, each spin is a function

$$\delta_i(\omega) = \begin{cases} 1 & \text{if } \omega_i \text{ is up} \\ -1 & \text{if } \omega_i \text{ is down} \end{cases} \quad (1.1)$$

An *energy* $U(\omega)$ is assigned to each configuration:

$$U(\omega) = -J \sum_{i,j} \delta_i(\omega) \delta_j(\omega) - mH \sum_i \delta_i(\omega). \quad (1.2)$$

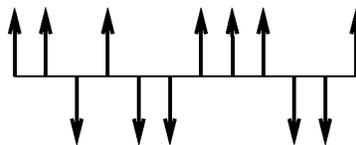


Fig. 1.1 One dimensional Ising model.

In the first sum, Ising made a simplifying assumption that only interactions of points with one unit apart need to be taken into account. This term represents the energy caused by the spin-interactions. The constant J is a property of the material. If $J > 0$, the interactions tend to keep neighboring spins in the same directions (*attractive case*). If $J < 0$, neighboring spins with opposite orientation are favored (*repulsive case*). The second term represents the influence of an external magnetic field of intensity H and $m > 0$ is a property of the material. The probability on Ω is then given by

$$P(\omega) = \frac{\exp\left(-\frac{1}{kT}U(\omega)\right)}{Z}, \quad (1.3)$$

where T is the temperature and k is a universal constant. The normalizing constant (also called *partition function*) Z is defined by

$$Z = \sum_{\omega \in \Omega} \exp\left(-\frac{1}{kT}U(\omega)\right). \quad (1.4)$$

The probability defined in Equation (1.3) is called a *Gibbs distribution*. One could extend the model to two dimensions in a natural way. The spins are arranged on a lattice, they are represented by two coordinates and a point has 4 neighbors unless it is on the boundary. In the two-dimensional case, the limiting measure P is unstable, there is a *phase transition*. As it is pointed out in [125], considering the *attractive case* and an external field h , the measure P_h converges to P^- if h goes to zero through negative values but it converges to $P^+ \neq P^-$ if h goes to zero through positive values. It has been shown, that there exists a *critical temperature* T_C and below this temperature phase transition always occurs. The temperature depends on the vertical (J_1) and horizontal (J_2) interaction parameters.

As a special example, we mention the *Cayley tree model* [125], originally proposed by Bethe [10] as an approximation to the Ising model. In this case, the points sit on a tree (see Figure 1.2). The root is called the 0th level. From the root, we have q branches ($q = 2$ in Figure 1.2). The $q = 1$ case simply gives a one-dimensional Markov chain. A configuration on a tree of n levels is an assignment of a label *up* or *down* to each point. We can define a similar energy function as for the Ising model.

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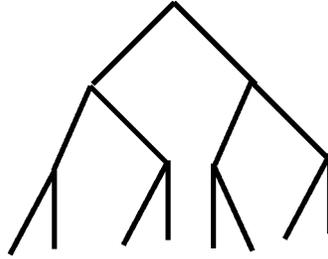


Fig. 1.2 Cayley tree model.

1.2.2 The Potts Model

Another important extension of the Ising model to more than two states per points is the Potts model [10, 195, 216]. The problem is to regard the Ising model as a system of interacting spins that can be either parallel or antiparallel. More generally, we consider a system of spins, each spin pointing one of the q equally spaced directions. These vectors are the linear combinations of q unit vectors pointing in the q symmetric directions of a hypertetrahedron in $q - 1$ dimensions. For $q = 2, 3, 4$, examples are shown in Figure 1.3. The energy function of the Potts model can be written as

$$U(\omega) = \sum_{i,j} J(\Theta_{ij}), \tag{1.5}$$

where $J(\Theta)$ is 2π periodic and Θ_{ij} is the angle between two neighboring spins in i and j . The $q = 2$ case is equivalent to the Ising model.

1.2.3 Gibbs Distribution and MRFs

The most natural way to define MRFs [2, 64, 184] related to image models is to define them on a lattice. However, here we will define

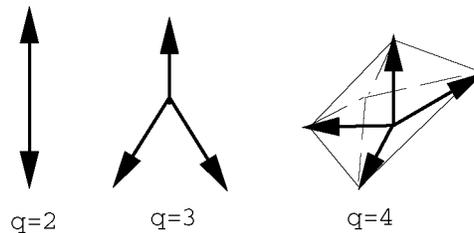


Fig. 1.3 The Potts model.

MRFs more generally on graphs. It will be useful in Section 2 for the study of hierarchical models. Let $\mathcal{G} = (\mathcal{S}, \mathcal{E})$ be a graph where $\mathcal{S} = \{s_1, s_2, \dots, s_N\}$ is a set of vertices (or sites) and \mathcal{E} is the set of edges.

Definition 1.1 (Neighbors). Two points s_i and s_j are neighbors if there is an edge $e_{ij} \in \mathcal{E}$ connecting them. The set of points which are neighbors of a site s (that is, the neighborhood of s) is denoted by \mathcal{G}_s .

Definition 1.2 (Neighborhood system). $\mathcal{G} = \{\mathcal{G}_s \mid s \in \mathcal{S}\}$ is a neighborhood system for \mathcal{G} if

- (1) $s \notin \mathcal{G}_s$
 - (2) $s \in \mathcal{G}_r \Leftrightarrow r \in \mathcal{G}_s$
-

To each site of the graph, we assign a label λ from a finite set of labels Λ . Such an assignment is called a configuration ω having some probability $P(\omega)$. The restriction to a subset $\mathcal{T} \subset \mathcal{S}$ is denoted by $\omega_{\mathcal{T}}$ and $\omega_s \in \Lambda$ denotes the label given to the site s . In the following, we are interested in the probabilities assigned to the set Ω of all possible configurations. First, let us define the *local characteristics* as the conditional probabilities $P(\omega_s \mid \omega_r, r \neq s)$.

Definition 1.3 (Markov random field). \mathcal{X} is a Markov random field (MRF) with respect to \mathcal{G} if

- (1) for all $\omega \in \Omega$: $P(\mathcal{X} = \omega) > 0$,
 - (2) for every $s \in \mathcal{S}$ and $\omega \in \Omega$:

$$P(X_s = \omega_s \mid X_r = \omega_r, r \neq s) = P(X_s = \omega_s \mid X_r = \omega_r, r \in \mathcal{G}_s).$$
-

To continue our discussion about probabilities on Ω , the notion of *cliques* will be very useful.

Definition 1.4 (Clique). A subset $C \subseteq \mathcal{S}$ is a clique if every pair of distinct sites in C are neighbors. \mathcal{C} denotes the set of cliques and $\text{deg}(\mathcal{C}) = \max_{C \in \mathcal{C}} |C|$.

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Using the above definition, we can define a *Gibbs measure* on Ω . Let V be a *potential* which assign a number $V_{\mathcal{T}}(\omega)$ to each subconfiguration $\omega_{\mathcal{T}}$. V defines an *energy* $U(\omega)$ on Ω by

$$U(\omega) = - \sum_{\mathcal{T}} V_{\mathcal{T}}(\omega). \quad (1.6)$$

Definition 1.5 (Gibbs distribution). A Gibbs distribution is a probability measure π on Ω with the following representation:

$$\pi(\omega) = \frac{1}{Z} \exp(-U(\omega)), \quad (1.7)$$

where Z is the normalizing constant (also called *partition function*):

$$Z = \sum_{\omega} \exp(-U(\omega)),$$

If $V_{\mathcal{T}}(\omega) = 0$ whenever \mathcal{T} is not a clique then V is called a *nearest neighbor Gibbs potential*. In the following, we will focus on such potentials. The next famous theorem establish the equivalence between Gibbs measures and MRFs [16, 160].

Theorem 1.6 (Hammersley–Clifford). \mathcal{X} is a MRF with respect to the neighborhood system \mathcal{G} if and only if $\pi(\omega) = P(\mathcal{X} = \omega)$ is a Gibbs distribution with a nearest neighbor Gibbs potential V , that is

$$\pi(\omega) = \frac{1}{Z} \exp \left(- \sum_{C \in \mathcal{C}} V_C(\omega) \right) \quad (1.8)$$

The main benefit of this equivalence is that it provides us a simple way to specify MRFs, namely specifying potentials instead of local characteristics (see Definition 1.3), which is usually very difficult.

1.2.4 Spatial Lattice Schemes

In this section, we deal with a particular subclass of MRFs which are the most commonly used schemes in image processing. In this case,

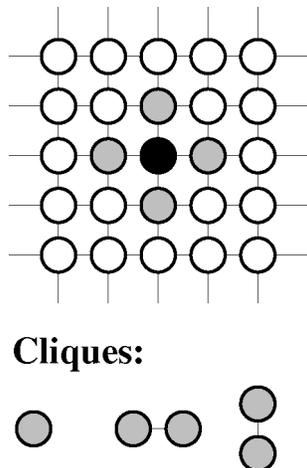


Fig. 1.4 First order neighborhood system.

we consider \mathcal{S} as a lattice \mathcal{L} so that $\forall s \in \mathcal{S} : s = (i, j)$ and define the so-called *n*th order homogeneous neighborhood systems as

$$\mathcal{G}^n = \{\mathcal{G}_{(i,j)}^n : (i, j) \in \mathcal{L}\}, \tag{1.9}$$

$$\mathcal{G}_{(i,j)}^n = \{(k, l) \in \mathcal{L} : (k - i)^2 + (l - j)^2 \leq n\}. \tag{1.10}$$

Obviously, sites near the boundary have fewer neighbors than interior ones (free boundary condition). Furthermore, $\mathcal{G}^0 \equiv \mathcal{S}$ and for all $n \geq 0 : \mathcal{G}^n \subset \mathcal{G}^{n+1}$. Figure 1.4 shows a first-order neighborhood corresponding to $n = 1$. The cliques are $\{(i, j)\}, \{(i, j), (i, j + 1)\}, \{(i, j), (i + 1, j)\}$. In practice, more than two order systems (cf. Figure 1.5) are rarely used since the energy function would be too complicated requiring a lot of computation. Although not as widespread as orthogonal lattice schemes, hexagonal lattices [45, 193] as well as MRFs on graphs [204] have also been studied in the literature.

1.3 Related Approaches

1.3.1 Weak Membrane Model

The *weak membrane model* was introduced in image reconstruction by A. Blake and A. Zisserman [19]. The problem is to reconstruct surfaces

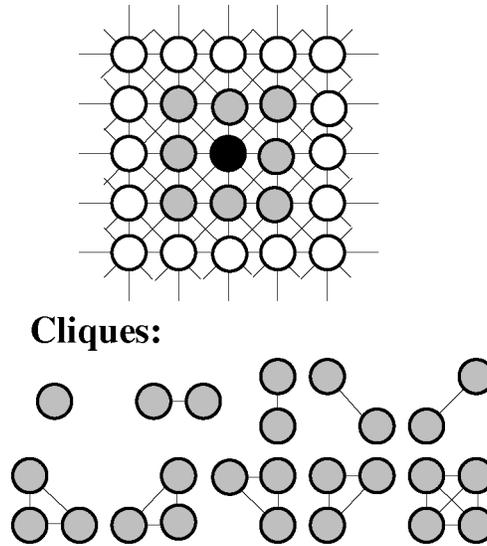


Fig. 1.5 Second order neighborhood system.

which are *continuous almost everywhere* or, in other words, continuous in patches. To reach a satisfactory formalization of this principle, they have used a membrane model: Imagine an elastic membrane which we are trying to fit to a surface. The edges will appear as tears in the membrane. Depending on how elastic is the membrane, there may be more or fewer edges. The membrane is described by an energy function (the elastic energy of the membrane) which has to be minimized in order to find an equilibrium state. The energy has three components:

D: A measure of faithfulness to the data:

$$D = \int (u - d)^2 dA, \quad (1.11)$$

where $u(x, y)$ represents the membrane and $d(x, y)$ represents the data.

S: A measure of how the function $u(x, y)$ is deformed:

$$S = \lambda^2 \int (\nabla u)^2 dA. \quad (1.12)$$

λ^2 is a measure of elasticity of the membrane.

P: The sum of penalties α levied for each break in the membrane:

$$P = \alpha Z, \quad (1.13)$$

where Z is a measure of the set of contours along which $u(x, y)$ is discontinuous (see [19] for more details).

The elastic energy of the membrane is then given by

$$E = D + S + P = \int (u - d)^2 dA + \lambda^2 \int (\nabla u)^2 dA + \alpha Z. \quad (1.14)$$

There is a strong relation between the *weak membrane model* and MRF models: An elastic system can also be considered from a probabilistic view-point. The link between the elastic energy E and probability P is

$$P \propto \exp\left(\frac{-E}{T}\right), \quad (1.15)$$

that is the Gibbs distribution. However, the *weak membrane model* operates with mechanical analogies, representing *a priori* knowledge from a mechanical point of view while MRF modelization is purely probabilistic.¹

1.3.2 Snakes, Variational and Level Set Methods

Active Contours (snakes) are closed curves evolving toward the boundary of the object of interest. The curve evolution is governed by a boundary functional [101] which takes its minimum on the object contour. The main drawback of the parametric snake model is that it cannot handle topological changes easily. Nevertheless, they became quite popular because they make it relatively easy to enforce contour-smoothness; and starting from an appropriate initialization a local minimum of the associated energy function will give good results. One extension of the original model is gradient vector flow [217] snakes

¹We notice that the weak membrane model has also been used in a Markovian context but originally, as proposed by Blake and Zisserman [19], it was a non-Markovian model.

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which make the snake less sensitive to initialization and allow the contour to segment concave objects. Another extension is the so-called balloon force [44] which basically introduces an area minimizing term [31] into the snake energy.

Geodesic active contours [31] are curves of minimum length in the metric defined by a function u . The criterion to minimize is usually of the form $\int_{\Gamma} u(s) ds$. Most of the time, u is simply a function of the image gradient like $u = 1/(1 + |\nabla \mathcal{I}|)$. The contour evolution equation is as follows [31]:

$$\frac{\partial \Gamma}{\partial t} = (\kappa u \nabla u \cdot N) N, \quad (1.16)$$

where κ is the curvature and N is the inward normal of Γ .

Region based active contours are another class of boundary based methods where region descriptors (usually some kind of statistical features) are introduced into the energy in order to better characterize an object [169, 188, 224].

Variational approaches consider the segmentation as an optimal approximation of the original image \mathcal{I} by a piecewise smooth function f having discontinuities across Γ . The classical Mumford–Shah energy functional [163] is then defined as

$$E(f, \Gamma) = \mu^2 \int \int_R (f - \mathcal{I})^2 dx dy + \int \int_{R-\Gamma} \|\nabla f\|^2 dx dy + \nu |\Gamma|. \quad (1.17)$$

Clearly, the minimum is achieved when f approximates \mathcal{I} (first term), f is smooth over each R_i (second term), and the boundaries Γ are as short as possible. Note that dropping any of the above three terms would result in $\inf E = 0$ with some trivial and (from a practical point of view) useless settings for f and Γ . The minimization of the above functional is far from trivial. Note also that in our context, $f = \omega$ is constant over each region R_i , hence the problem can be further simplified to a piecewise constant functional. A closely related model, proposed by Blake and Zisserman, is the so-called *weak membrane* model (see Section 1.3.1) which can be minimized via graduated non-convexity (GNC) [19].

More recently, the level set formulation [192] of the piecewise constant Mumford–Shah energy functional proposed by Chan and Vese [38] have become a popular framework for image segmentation. The contour Γ is represented as the zero level set of an embedding function (the level set function) $\phi : R \rightarrow \Re$ on the image domain R : $\phi(\Gamma) = 0$. The main advantage of this formulation is that it handles topological changes of the evolving contour. This makes the level set formalism well suited to the segmentation of multiple objects. The region based level set scheme for foreground–background segmentation consists in minimizing the following functional:

$$E_{CV}(c_1, c_2, \phi) = \int_R (\mathcal{I} - c_1)^2 H(\phi) dx + \int_R (\mathcal{I} - c_2)^2 (1 - H(\phi)) dx + \nu \int_R |\nabla H(\phi)| dx, \quad (1.18)$$

c_1 and c_2 are the means of the regions, where $\phi > 0$ (outside or background) and $\phi < 0$ (inside or foreground), and $H(\cdot)$ is the Heaviside function. The last term measures the length of the zero crossing of ϕ (i.e., the contour). The Euler–Lagrange equation for this model is implemented by the following gradient descent:

$$\frac{\partial \phi}{\partial t} = \delta(\phi) \left[\nu \operatorname{div} \left(\frac{\nabla \phi}{|\nabla \phi|} \right) - (\mathcal{I} - c_1)^2 + (\mathcal{I} - c_2)^2 \right]. \quad (1.19)$$

Unfortunately, even with the narrow band implementation [1], the level set approach has a rather high computational complexity. The fast marching method [192] has a lower complexity but it requires that the speed function doesn't change sign during evolution.

1.3.3 Conditional Random Fields

Conditional Random Fields (CRF) directly model the posterior distribution of $P(X|Y)$ as a Gibbs field [86, 135, 136, 206]. Unlike the generative image models commonly used in MRFs, CRFs can depend on arbitrary non-independent characteristics of the observation $Y = y$. Originally, CRFs were proposed for segmenting 1D text

sequences [140, 212], but it is straightforward to extend these concepts to 2D images.

Basically, a CRF is a random field globally conditioned on the observation Y . Following [140], we can formally define CRFs on graph:

Definition 1.7 (Conditional Random Field). Let $G = (V, E)$ be a graph such that the label field X is indexed by the vertices: $X = \{X_v\}_{v \in V}$ and neighboring elements $v \sim w$ of the field are connected by edges in G , i.e., $(v, w) \in E$. Then (Y, X) is a *conditional random field (CRF)* if the random variables X_v , when conditioned on Y , obey the Markov property with respect to the graph: $P(X_v|Y, X_w, w \neq v) = P(X_v|Y, X_w, w\tilde{v})$.

The simplest example of such a graph structure is a lattice where vertices correspond to pixels and neighboring lattice sites are connected by edges (see Section 1.2.4 for various neighborhood structures on lattices). Considering a first order neighborhood, the posterior distribution can be easily expressed using the Hammersley–Clifford theorem (see Theorem 1.6):

$$P(x|y) = \exp \left(\sum_{e \in E, k} \lambda_k f_k(e, x|_e, y) + \sum_{v \in V, k} \mu_k g_k(v, x|_v, y) \right), \quad (1.20)$$

where x is a labeling of a given input image y and $x|_S$ is the set of components of x associated to the vertices in the subgraph S . Furthermore, the features f_k and g_k are assumed to be known and fixed, and the parameter values λ_k and μ_k are to be learned from training data [140]. As we can see from Equation (1.20), standard CRFs use two forms of feature functions, which can be interpreted in 2D as follows [86]:

- state feature function $g_k(s, x_s, y)$ of the label x_s at a site s and the observed image y ,
- transition feature function $f_k(s, r, x_s, x_r, y)$ of the labels x_s and x_r at neighboring sites $s \sim r$ and the observed image y .

In image processing applications, state feature functions are usually defined as unary (also known as singleton) clique potentials based

on classifier responses (such as Ada-boost [194] or kernel SVMs [191]), while transition feature functions are defined as pairwise (also known as doubleton) potentials modeling the correlation between pairs of random variables. Recently, CRFs became popular in image segmentation [86, 206], especially CRFs coupled with graph cut energy minimization [128, 138, 208].

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