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Content-based image retrieval tools and techniques

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Content-Based Image Retrieval Tools and Techniques



In the beginning was an image.

To my mother who inspired me to develop intellectually

]



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3 Image Representations

3.1 Introduction. Forms of Image Representation

With regard to computers, an image can be represented in different forms. The two most frequently used types use at present are vector and raster images. Vector graphics, generated as geometric shapes by mathematical equations, can be scaled, rotated, moved, or otherwise manipulated to any degree without any loss of quality, and displayed or printed at whatever resolution is available on a monitor or printer. An example of a vector image is shown in Fig. 3.1 a) as a line image and in Fig. 3.1 b) as a colour one. In turn, raster graphics is made up of pixels, each of a different colour, arranged to display an image. A major difference is that raster image pixels do not retain their appearance as size increases, when you blow a photograph up, it becomes blurry for this reason. These two representations can be compared in Fig. 3.1 c) vector zoom and d) raster zoom.

Here, we analyse visual information so, first of all, we have to explain how we understand the notion of image.

Definition 3.1 (image)

An image is a two dimensional function $f : \mathbb{R}^2 \to \mathbb{R}$ such as f(x,y) describing the intensity at the position (x,y). For digital images *I*, we have a pixel array *I*: $[a,b] \times [c,d] \to [0,1]$. Colour images consist of three colour components:

$$I: (x,y) = \begin{bmatrix} r(x,y) \\ g(x,y) \\ b(x,y) \end{bmatrix}$$
(3.1)

Generally, we analyse and store in the DB raster images, but in some cases we use vector images, for example, as a prompt for the user in the GUI (see Fig. 3.2), where we exploit the easy zooming (details in Chapter 8).

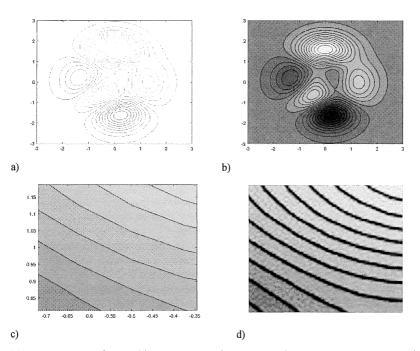


Fig. 3.1 Two most often used image representations: raster and vector; a) vector representation, b) vector with colour filling, c) close-up of the vector representation, d) close-up of the raster representation.

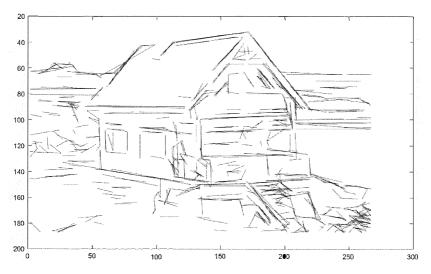


Fig. 3.2 Example of a vector image - used as a prompt in the GUI.

3.2 Visual Feature Descriptors

Feature extraction is a process of selecting a map of the form X=f(Y), by which a sample $\mathbf{y}=[y_1,y_2,...,y_q]$ in a *q*-dimensional measurement space Ω_Y is transformed into a point $\mathbf{x}=[x_1, x_2, ..., x_{q'}]$ in a *q*'-dimensional feature space Ω_X , where q' < q. This task is realized to generate the optimal characteristics necessary for the process of recognition and reduce the dimensionality of space Ω_Y in order to apply effective computable algorithms of classification. We understand the space Ω_Y as a model which we construct based on a subset of selected variables.

The exact definition of feature often depends on the problem or the type of application. Feature detection is low-level image processing because it operates on the level of pixels and is the easiest to extract by the computer system. As a matter of fact, all CBIR systems use the feature detection level, some concentrate on one feature, for instance, colour or shape, others use their combination, but only the more advanced move further to semantic retrieval.

Below we present the most commonly used algorithms which are also useful in search engine construction. We would like to present a wide range of methods as a background for our content-based image retrieval system.

To begin with, we describe our approach to colour segmentation as the first point in constructing our CBIR system which, step by step, will be presented throughout this book.

3.3 Colour Information

Colour is a commonly used feature because its layout in the image is the key information, whereas the simpler systems extract only global features from the colour image. The more advanced ones use colour information about regions or separate segments [34].

Each pixel of the image can be represented as a point in a 3D colour space. Many colour spaces for image retrieval, including *RGB*, *Munsell*, *CIE* $L^*a^*b^*$, *CIE* $L^*u^*v^*$, *HSV* are used depending on the aims and the method of image acquisition.

Colour information gives the opportunity to construct such descriptors as:

- colour moments (mean, variance and skewness) [56], [17] help to describe colour distribution in the whole image, which is the basis for many CBIR retrieval processes. Nevertheless, they do not give spatial information about pixels;
- **the colour histogram** is easy to compute and invariant in terms of scaling and rotation, however, it also fails to provide spatial information about pixels, so many images have similar histograms;
- the colour coherence vector (CCV) [57] is constructed based on the colour histogram. In this case, each histogram bin, a separate one for each colour, is

partitioned into two parts: coherent if it belongs to a large uniformly-coloured region, and incoherent in the opposite case. It means that two pixels *a* and *a'* are coherent if they belong to region *C*, such that $a, a' \in C$ and there exists a path in *C* between *a* and *a'*. For the image, the CCV is defined as the vector $[(\alpha_1, \beta_1), (\alpha_2, \beta_2), ..., (\alpha_N, \beta_N)]$, where α_i denotes the number of coherent pixels of the *i*th colour bin, whereas β_N denotes the number of coherent pixels. The additional spatial information included in the CCV improves the results of retrieval in comparison to the simple colour histogram.

• the colour correlogram, also called a second-order histogram, describes the spatial correlation of pairs of colours. A colour correlogram is a table indexed by colour pairs, where the k^{th} entry for (i,j) specifies the probability of finding a pixel of colour j at a distance r from a pixel of colour i in the image. The colour correlograms for all sets of colours are rather large, therefore, a simplified version is an auto-correlogram which is a spatial relationship only between points of identical colour.

In our approach, we omitted the colour description globally. We only used colour information, specifically RGB colour space, to segment separate objects from an image (in details see sec. 4.2).

3.3 Texture Information

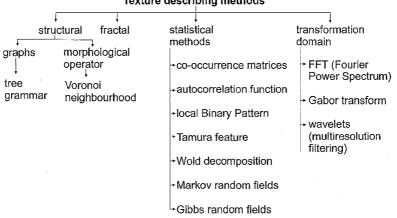
Texture is one of the most important visual cues to identify homogeneous regions [58]. The goal of texture classification is to identify each uniform texture region, whereas the goal of texture segmentation is to obtain the boundary map and further separate regions characterized by different textures.

For our purpose, the key operation is segmentation, in a nutshell, we can assume that image texture is an attribute representing the spatial arrangement of grey or colour levels of the pixels in a region [59]. Hence, the intensity variations in an image, characterizing texture, generally reflect physical variations in the real scene. To model these variations the following issues need to be addressed:

- pixel colour value in a spatial neighbourhood;
- spatial distributions of these values;
- their resolution or scale;
- the unrecognizability of separate primitive objects in a texture region.

Basically, texture representation methods can be classified into four categories: structural, statistical, fractal [60] and transformational [34] as it can be seen in Fig. 3.. The first category of methods can be divided into morphological operators and adjacency graphs presenting texture as structural primitives and their placement rules. The primitive can be as simple as a single pixel that can take a grey value, but it is usually a collection of pixels. The placement rule is defined by a tree grammar. A texture is then viewed as a string in the language defined by the

grammar whose terminal symbols are the texture primitives. An advantage of this method is that it can be used for texture generation, as well as texture analysis. The patterns generated by the tree grammars can also be regarded as ideal textures in Zucker's model [61]. They are more effective when we have a regular texture.



Texture describing methods

Fig. 3.3 The categories of texture describing methods.

Another example is Voronoi features [62], which were proposed because the local spatial distributions of tokens are reflected in the shapes of Voronoi polygons. Many of the perceptually significant characteristics of a token's environment are manifest in the geometric properties of Voronoi neighbourhoods. In order to apply geometrical methods to grey level images, we only need to first extract tokens from images.

The statistical methods describe texture by statistical distribution of image intensity. There are numerous statistical texture representations:

co-occurrence matrices. Spatial grey level co-occurrence estimates image properties related to second-order statistics. Haralick [63] suggested the use of the $G \times G$ grey level co-occurrence matrix P_d for a displacement vector $\mathbf{d} = (dx, dy)$, defined as follows: the entry (i, j) of $P_{\mathbf{d}}$ is the number of occurrences of the pair of grey levels i and j which are a distance **d** apart. Formally, it is given as:

$$P_{\mathbf{d}}(i,j) = |\{((x,y), (t,v)) : I(x,y) = i, I(t,v) = j\}|$$
(3.2)

where $(x,y),(t,v) \in N \times N$, (t,v) = (x+dx, y+dy), and |.| is the cardinality of a set. Based on this matrix some useful texture features can be described, such as: energy, entropy, contrast, homogeneity or correlation.

Autocorrelation function - can be used to assess the amount of regularity as well as the fineness – coarseness of the texture in the image. Formally, the autocorrelation function of an image I(x,y) is defined as follows:

$$\sigma(x,y) = \frac{\sum_{u=0}^{N} \sum_{\nu=0}^{N} I(u,\nu) I(u+x,\nu+y)}{\sum_{u=0}^{N} \sum_{\nu=0}^{N} I^{2}(u,\nu)}$$
(3.3)

Local Binary Pattern (LBP) operator [64] is, originally, based on a 3×3 pixel neighbourhood (see the *example* sub-table in Table 3.1). Image pixels in each neighbourhood of a pixel (i,j) are exchanged into a binary threshold map where 1 is for a pixel larger than the central pixel and 0 where the values are less than the central one (see the *threshold* sub-table in Table 3.1). The values of the pixels in the threshold map are multiplied by the weights given to the corresponding pixels. The weights are the power of 2 where the number of neighbourhood is the exponent (see the weights sub-table in Table 3.1). Finally, the values of the eight weighted pixels are summed to obtain one factor (the result can be seen in Table 3.1). The LBP histogram computed over a region is used as a texture description. Because of the LBP design, it is invariant under any monotonic grey scale transformation and provides information about the spatial structure of the local image texture. Due to its 3×3 window operation, however, feature distributions may be sensitive to geometric distortion. This operator was extended later [65] for neighbourhoods of different sizes, for instance, circular neighbourhood and bilinear interpolation of non-integer pixel values.

E	xamp	le		Th	resh	old			Weig	hts]	Result	t
8	5	2		1	0	0		1	2	4	1	0	0
7	6	1		1	x	0		8	x	16	8	x	0
9	13	7		1	1	1		32	64	128	32	64	128
I BP = 1 + 8 + 32 + 64 + 128 = 233													

 Table 3.1 Computation of Local Binary Pattern (LBP).

LBP = 1 + 8 + 32 + 64 + 128 = 233

The original LBP operator was defined to only deal with the spatial information. Later, it was extended to a spatiotemporal representation for dynamic texture analysis. For this purpose, the so-called Volume Local Binary Pattern (VLBP) operator was proposed [66].

Tamura feature; Tamura et al. [67] considered six basic textural features:

- coarseness relates to distances of notable spatial variations of grey levels, that is, implicitly, to the size of the primitive elements forming the texture.
- contrast measures how grey levels vary in the image and to what extent their distribution is biased to black or white. The second-order and normalised fourth-order central moments of the grey level histogram are used to define the contrast.
- directionality measured the distribution of oriented local edges against their directional angles using the Sobel edge detector (for details see sec. 3.4.1).
- line-likeness is defined as an average coincidence of the edge directions.

- regularity is defined as the normalised sum of the standard deviations of the corresponding above-mentioned feature.
- roughness feature is given by simply summing the coarseness and contrast measures.
- Wold decomposition, [68], [59] provides three different components to describe texture: *harmonic, evanescent*, and *non-deterministic*, corresponding to *periodicity, directionality*, and *randomness* introduced by his predecessors. Periodic textures have a strong harmonic component, highly directional textures have a strong evanescent component, and less structured textures tend to have a stronger non-deterministic component. The deterministic periodicity of the image is analysed using the autocorrelation function. The corresponding Wold feature set consists of the frequencies and the magnitudes of harmonic spectral peaks (e.g. the largest peaks). The nondeterministic (random) components of the image are modelled with the multiresolution simultaneous autoregressive (MR-SAR) process. The retrieval uses matching of the harmonic peaks and the distances between the MR-SAR parameters. The similarity measure involves a weighted ordering based on the confidence level in the query pattern regularity.
- Markov random fields [69]. Random field models consider an image as a 2D array of random scalars (grey values) or vectors (colours). In other words, the signal at each pixel location is a random variable. Each type of texture is characterised by a joint probability distribution of signals that accounts for spatial inter-dependence, or interaction among the signals. The interacting pixel pairs are usually called neighbours, and a random field texture model is characterized by the geometric structure and quantitative strength of interactions among the neighbours. If pixel interactions are assumed translation invariant, the interaction structure is given by a set N of characteristic neighbours of each pixel. This results in the Markov random field model where the conditional probability of signals in each pixel (i,j) depends only on the signals in the neighbourhood $\{(i+m, j+n): (m,n)$ from the set $N\}$.
- Gibbs random fields (GRF). This theory is borrowed from Gibbs principal ensembles of statistical thermodynamics. We move from particles to pixels and still analyse potential and energy functions. Hence, GRF assigns a probability mass function to the entire lattice:

$$P(\mathbf{X} = \mathbf{x}) = \frac{1}{Z} \exp\left[-\sum_{c_i \in C} E(c_i)\right], \quad \forall \mathbf{x} \in \Omega$$
(3.4)

where Z is a normalizing constant known as the partition function and $E(c_i)$ is the energy function.

For texture analysis, general generic Gibbs random field models with multiple pairwise pixel interactions allow to relate the desired neighbourhood to a set of most 'energetic' pairs of the neighbours. A Gibbs distribution is usually defined with respect to cliques, i.e. proper subgraphs of a neighbourhood graph on the lattice. A clique is a particular spatial configuration of pixels, in which all its members are statistically dependent on each other. Then the interaction structure itself and relative frequency distributions of signal co-occurrences in the selected pixel pairs can serve as texture features.

Many natural surfaces have a statistical quality of roughness and self-similarity at different scales. Fractals are very useful and have become popular in modelling these properties in image processing but scale variations can have a great impact on the imaged appearance of a texture. Self-similarity across scales in fractal geometry is a crucial concept. The fractal dimension gives a measure of the roughness of a surface.

Fractal-based texture analysis was introduced by Pentland in 1984 [70]. To apply the fractal model to an image surface, we need to assume that: the intensity random function I(x) is a fractal Brownian function and the fractal dimension of a fractal Brownian function is invariant over transformations of scale¹. In order to obtain the fractalness of an image, Pentland introduced the description of the image change $\Delta I = I(x + \Delta x) - I(x)$ with scale as follows (eq. (5) [70]):

$$E(|\Delta I_{\Delta x}|) ||\Delta x||^{-H} = E(|\Delta I_{\Delta x=1}|)$$
(3.5)

where: $E(|\Delta I_{\Delta x}|)$ is the expected value of the change in intensity ΔI over distance Δx , *H* is the Hurst exponent [71], [72]. Equation (3.5) is the mutual relation of the image intensities expressed in a statistical way.

We can assume that $\kappa = E(|\Delta I_{\Delta x}=1|)$, hence we obtain in the above equation $E(|\Delta I|) = \kappa ||\Delta x||^{H}$. By applying log to both sides we have

$$\log E(|\Delta I|) = \log \kappa + H \log ||\Delta x||.$$
(3.6)

The Hurst exponent H can be obtained by using the linear least-squares regression to estimate the slope of the grey-level difference GD(k) versus k in a log-log scale over the interval k = [1,s], s - the maximum value, where:

¹ Following Mandelbrot [276] the increments of a random function $\{X(t,\omega); -\infty \le t \le \infty\}$ are said to be self-similar with parameter $H \ge 0$ if for any h > 0 and any moment t_0

$$\{X(t_0+\tau,\omega) - X(t_0,\omega)\} \triangleq \{h^{-H}[X(t_0+h\tau,\omega) - X(t_0,\omega)]\}.$$

If $X(t_0, \omega)$ has self-similarity and stationary increments and is mean square continuous, then $0 \le H \le 1$ there is a constant V such that

$$E\left[X(t_0+\tau,\omega)-X(t_0,\omega)\right]^2=V\tau^{2H}.$$

For images, following Pentland [70], instead of time t we speak about spatial dimension x, so we have $E[I(x+\Delta x) - I(x)]^2 = V \Delta x^{2H}$.

$$GD(k) = \frac{\sum_{x=1}^{N} \sum_{y=1}^{N-k-1} |I(x,y) - I(x,y+k)|}{2N(N-k-1)} + \frac{\sum_{x=1}^{N-k-1} \sum_{y=1}^{N} |I(x,y) - I(x+k,y)|}{2N(N-k-1)}$$
(3.7)

The fractal dimension FD can be derived from the relation FD=3-H. The approximation error of the regression line fit should be determined to prove that the analyzed texture is fractal, and thus be efficiently described using fractal measures. A small value of the fractal dimension FD implies that a large value of the Hurst exponent H represents fine texture, while a large FD, implying a smaller H value, corresponds to coarse texture [73].

Recently, texture descriptors have been based on transformational models. Let us recall the basic notions of the unitary transform. A general linear operation on the input image I(x,y) results in an $M \times N$ output image U(m,n) which is defined by:

$$U(m,n) = \sum_{x=0}^{K-1} \sum_{y=0}^{J-1} I(x,y) O(x,y;m,n)$$
(3.8)

where: O(x, y; m, n) is the operator kernel.

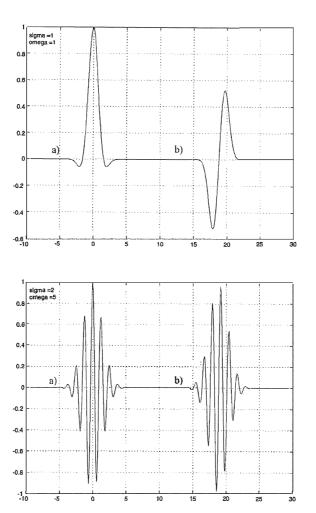
Based on this universal rule we can chronologically describe the most useful and therefore the most common transformational methods:

• Fourier power spectra and Fast Fourier Transform (FFT) [74]. For image function I(x, y) we compute its Fourier transform as:

$$F(u,v) = \frac{1}{N} \sum_{x=0}^{N-1} \sum_{y=0}^{N-1} I(x,y) \exp\left\{\frac{-2\pi i}{N} (xu+yv)\right\}$$
(3.9)

where (u,v) are the spatial frequencies and the quantity $|F(u,v)|^2$ is defined as the power spectrum which, in fact, is the modulus of a complex number. In the image terms the energy distribution of the power spectrum reflects the periodical structure of a texture, whereas the directional nature of the texture is reflected in the direction distribution of energy in the power spectrum. Frankly speaking, the limitation at high frequencies is the image resolution.

• The Gabor transform [75], [76]. The Fourier transformation is an analysis of the global frequency content in the signal. Many applications require the analysis to be localized in the spatial domain. This is usually handled by introducing spatial dependency into the Fourier analysis. The classical way is using the windowed Fourier transform. Considering one dimension, the windowed Fourier transformation of a sinusoidal wave $f_{u_0}(x) = e^{iu_0x}$ (illustrates in Fig. 3.4) is defined as: $F(u) = 2\pi\delta(u - u_0)$, where δ is the Dirac



$$\left[u_0 - \frac{\sigma_u}{2}, u_0 + \frac{\sigma_u}{2}\right].$$

Fig. 3.4 Gabor function, where a) the real part of the function and b) the imaginary part of the function [79].

When the 2D window function w(r) is Gaussian,

$$w(r) = \frac{1}{2\pi\sigma^2} e^{-\frac{r^2}{2\sigma^2}}; \qquad r^2 = x^2 + y^2$$
(3.10)

the transform becomes a 2D Gabor transform [77], [78]:

$$G(x,y) = \frac{1}{2\pi\sigma_x\sigma_y} \exp\left\{-\pi \left[\frac{(x-x_0)^2}{\sigma_x^2} + \frac{(y-y_0)^2}{\sigma_y^2}\right]\right\} \cdot e^{i(u_0x+v_0y)}$$
(3.11)

where (x_0, y_0) is the centre of the receptive window in the spatial domain and (u_0, v_0) is the optimal spatial frequency of the filter in the frequency domain. σ_x and σ_y are the standard deviations of the elliptical Gaussian along x and y. The 2D Gabor function is thus a product of an elliptical Gaussian and a complex plane wave.

The 2D Gabor function consists of a sinusoidal plane wave of a certain frequency and orientation modulated by a Gaussian envelope given by:

$$g(x,y) = \exp\left\{-\frac{1}{2}\left[\frac{x^2}{\sigma_x^2} + \frac{y^2}{\sigma_y^2}\right]\right\} \cos(2\pi\omega_0(x\cos\theta + y\sin\theta))$$
(3.12)

where ω_0 and θ are the frequency and phase of the sinusoidal wave, respectively. Then a set of Gabor filters can be obtained by appropriate dilations and rotations of g(x,y) for angles $\theta = \frac{n\pi}{K}$, n = 0,1,...,K-1 (see Fig. 3.5). In this case, the Gabor transform of an image I(x,y) is defined as:

$$W_n(x,y) = \int I(x,y) \,\overline{g_n}(x - x_0, y - y_0) dx_0 dy_0 \tag{3.13}$$

for which:

$$\mu_n = \int |W_n(x, y)| \, dx \, dy \tag{3.14}$$

$$\sigma_n = \sqrt{\int (|W_n(x,y)| - \mu_n)^2 dx dy}$$
(3.15)

where μ_n is the mean and σ_n is the standard deviation of the magnitude of $W_n(x,y)$ for a particular orientation, where *n* denotes the specific subband. The texture analyzers based on 2D Gabor functions offer a strong correlation with the actual human segmentation and respective visual field profiles.

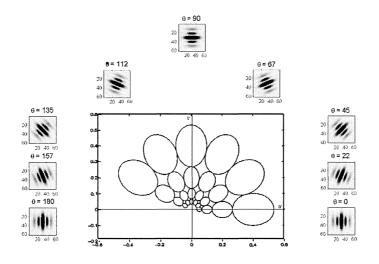


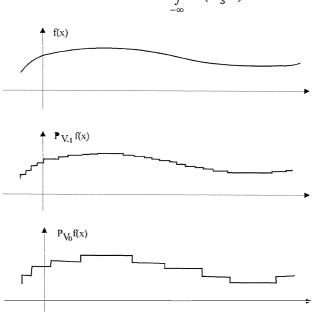
Fig. 3.5 Examples of 2D Gabor functions for particular angles $\theta = \frac{n\pi}{K}$, where K is the number of orientations. The outside windows present 2D Gabor filters, where K = 9. The central contours correspond to the half-peak magnitude of the filter responses in the set of Gabor filters with the upper and lower centre frequency of interest: $\omega_h = 0.4$ and $\omega_l = 0.05$, respectively, six orientations (K = 6), and four scales (S = 4), followed by [80].

• The wavelet transformations are a big group of methods focused on the multiresolution analysis concept. Generally, the structures to be recognized differ significantly in size. Hence, it is impossible to define *a priori* an optimal resolution for image analysis. Burt [81] and Crowley [82] have each introduced pyramidal implementation to compute image details in different resolutions. A multiresolution analysis (MRA) yields a scale-invariant interpretation of the image. A multiresolution representation provides a simple hierarchical framework for interpreting the image information. In different resolutions, details of an image generally characterize different physical structures of the scene; in a coarse resolution, these details correspond to larger structures represented by 'big' image components.

The idea of wavelets is based on a basic function called a wavelet (3.16) with two parameters: one - s, characterizing the scale, the other one - u, indicating the position of the function, introduced instead of the sinusoidal basic function with one parameter ω in the Fourier transform.

$$\psi_{su}(x) = \frac{1}{\sqrt{s}}\psi\left(\frac{x-u}{s}\right) \tag{3.16}$$

Hence, the 1D continuous wavelet transform is the projection of an f(x) signal, in the $L^2(R)$, onto the function family $\{\psi_{su}, s > 0, u \in R\}$ generated from the single function ψ by translation and dilation:



$$\left[W_{\psi}f\right](s,u) = \langle \psi_{su}, f \rangle = s^{-\frac{1}{2}} \int_{-\infty}^{+\infty} \overline{\psi\left(\frac{x-u}{s}\right)} f(x)dx$$
(3.17)

Fig. 3.6 A function f(x) and its projection onto two consecutive levels V_{-1} and V_0 of the multiresolution analysis [83].

The idea of this method is presented in Fig. 3.6.

The redundancy of the continuous wavelet transform (3.17) can be cleared by discretizing both the scale factor *s* and the translation *u*. Then, we need a dyadic scale space, $s=2^{j}$ and u=k with $j,k\in Z$ where Z is an integer. The fragment of the orthogonal basis with levels from V_{-3} to V_{-7} for Symmlet wavelets can be seen in Fig. 3.7.

The theory of multiresolution signal decomposition was developed by Mallat [84], [85] and Meyer [86] and thus the paradigm for constructing wavelets was established. Polish mathematicians were also involved in this analysis [87].

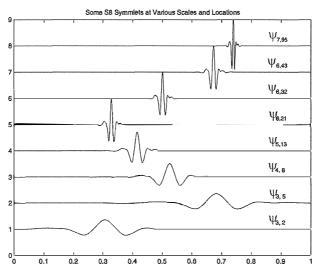


Fig. 3.7 An example of the dyadic Symmlet wavelets. A scale *j* and location *k* are presented for each wavelet $\psi_{j,k}$ on the right side [79].

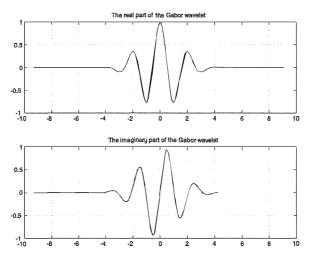


Fig. 3.8 The real and imaginary parts of the Gabor wavelet for $\sigma = 2$ and $\omega = 3$ which are 'larger' than the example of the subset shown in Fig. 3.5 [79].

Many different wavelets have been introduced over the years, some of them are real and some others are complex. The Gabor wavelet (see Fig. 3.8) is an example of a function in the complex domain. Each ellipse from Fig. 3.5 represents the frequency support of a dyadic wavelet $\hat{\psi}_{2j}^{K}$. This support size is proportional to 2^{j} and its position rotates when K is modified.

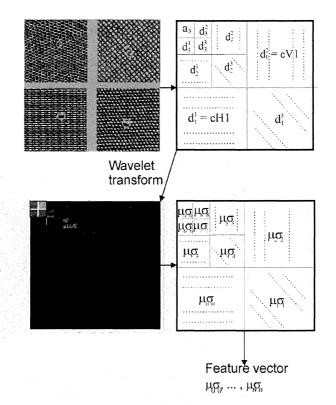


Fig. 3.9 A texture classifier flow chart based on the Gabor wavelet transformation (follows [80], [88]).

The discrete Gabor wavelet transform (GWT), for example, is applied in texture recognition and segmentation. Sebe and Lew [80] prepared an efficient method based on GWT parameters, such as μ and σ to be used as the texture feature vector. Fig. 3.9 presents four different grey texture representatives (top left) and the organization of wavelet image coefficients $d_{j,k,l}^p = \langle I, \psi_{j,k,l}^p \rangle$ [88]. The dotted lines show the direction of details (top right). In the bottom left square we can see a wavelet transform for texture images, whereas in the bottom right square we can see the organization of GWT coefficients (cf. (3.14) and (3.15)) that constitute feature vector $f = \{\mu_n, \sigma_n\}$.

At the same time, Faizal and Fausi [89] also used the DWT decomposition scheme and they noticed that the DWT of image $M \times M$ gave as a result $(3K+1) \times M^2$ coefficients. Based on this fact, they transformed wavelet coefficients to the 3D domain where they looked for clusters whose centroids characterized a texture.

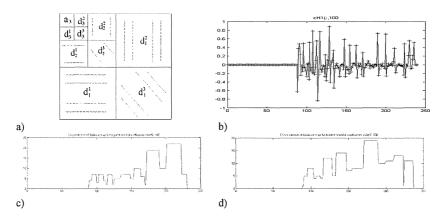
3.3.1 The Texture Approach to the Hybrid Semantic System

One of the reasons why so many kinds of wavelets were introduced in the previous section was to apply them to shape description. Our aim was to describe precisely the surface shape of an image function and we found out that the 'shape' of different wavelets corresponds more or less with the surface shape of the analysed image function. We should mention that from the shape matching point of view, one of the most important features of 2D wavelets is their detail directionality which is shown in Fig. 3.10 a) (called by Mallat as coefficients [88]). Thanks to this fact the convolution of wavelet and image element results in maximum values for concurrent element shapes (compare Fig. 3.9).

This wavelet property was applied by Jaworska [22], in order to describe the shape and size of the geometrical architectural texture. These parameters are different ones than those defined by Sebe and Lew or Faizal and Fausi in the previous subsection. We used the simplest wavelets, namely, the Haar wavelets which best fit a geometrical texture.

If we compute the convolution of an image consisting of regular tiles or bricks and vertical and horizontal details, we obtain a 2D transform whose maximum values are placed in the connection spots among these tiles or bricks. One dimension example of wavelet coefficients obtained from the convolution horizontal details and the 100^{th} line of image segment is shown in Fig. 3.10 b).

Having computed horizontal details, we have measured separately distances between maxima (shown in Fig. 3.10 c)) and between minima for each column of this matrix (shown in Fig. 3.10 d)). In this way, we created a 2D distance map in one direction for the analysed object. Repeating this procedure for vertical details, we obtained the second 2D distance map for the other direction. Based on 2D FWT maps of the object, texture is parametrised calculating the change distributions in the horizontal and vertical direction, respectively (see Fig. 3.10 g) and h)).



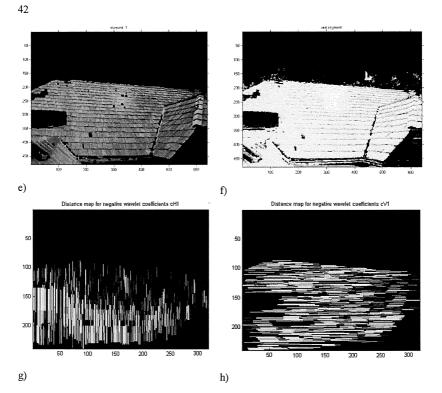


Fig. 3.10 Distance maps of texture calculated based on the 2D FWT with Haar wavelets. a) The disposition of wavelet image coefficients d_j^p where *j* is a multiresolution level, and a_j is an approximation at *j* level. b) Horizontal wavelet coefficients presented along the 100th column of the image transform (for the Haar wavelet, where j = 1). c) Cross-section through the 100th column of the distance map for positive horizontal wavelet coefficients. d) Cross-section through the 100th column of the distance map for negative horizontal wavelet coefficients. e) Original image of a roof segment (the segment was separated from the whole image based on our colour algorithm (cf. subsect. 4.2.3). f) The red component of the original image. g) Distance map for negative horizontal wavelet coefficients cV1 [49].

As a result of the above described algorithm, we obtain two ranges for the horizontal texture object components h and two others for the vertical one v.

$$T_{p} = \begin{bmatrix} h_{\min_{1,2}}; h_{\max_{1,2}} \\ [v_{\min_{1,2}}; v_{\max_{1,2}}] \end{bmatrix}$$
(3.18)

3.4 Edge Detection

We understand an edge as a discontinuity in the image brightness which helps us to identify curved lines – edges separated segments. Therefore, in an ideal case, the result of applying an edge detector to an image may lead to a set of connected curves that indicate the boundaries of objects, the boundaries of surface markings, as well as curves that correspond to discontinuities in surface orientation.

Generally, to detect edges we have to assume an image model in which discontinuities of image brightness are likely to correspond to:

- discontinuities in depth;
- discontinuities in surface orientation;
- · changes in material properties;
- variations in scene illumination.

The edge or contour can be defined as a parametric curve, polygon, or B-spline, but this can cause problems with the description of non-uniformed topological objects.

The method presented below covers grey images, because edge detection for colour images is more complicated. If a pixel falls on the boundary of an object in an image, then its neighbourhood is a zone of grey-level transition. Edge detection operators examine each pixel neighbourhood, and quantify the slope, as well as the direction, of the grey level transition.

There are several methods to do this, for example:

- watershed algorithm [91];
- gradient methods;
- active contours;
- Hough transform;
- fuzzy thresholding [92];

3.4.1 Gradient Methods

In gradient methods we treat the slope and direction of a potential edge as the magnitude and direction of the gradient vector, respectively, we apply the second derivative of the intensity of a 2D image function I(x,y), namely, the Laplacian:

$$\nabla^2 I(x,y) = \frac{\partial^2}{\partial x^2} I(x,y) + \frac{\partial^2}{\partial y^2} I(x,y)$$
(3.19)

thresholding [90]

The Laplacian is a linear, shift-invariant operator and its transfer function is equal to zero at the origin of a frequency space. Fig. 3.11 presents an example of edges and both derivatives.

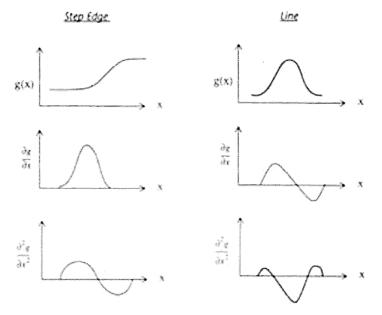


Fig. 3.11. The kind of edges (at the top), the first derivative of the edges (in the middle), the second derivative of the edges (at the bottom).

At present, for the discrete image version, most methods are based on convolution with a set of directional derivative masks M -filters:

$$\mathbf{H} = \mathbf{I} * \mathbf{M} \tag{3.20}$$

$$H(m,n) = \sum_{i} \sum_{j} I(i,j) M(m-i,n-j)$$
(3.21)

where the exemplary masks $\mathbf{M}_{[3\times 3]}$, for a discrete Laplacian H(m,n) are shown in Table 3.2.

Table 3.2 Laplacian convolution kernels.

The other well-known edge operators are suggested by: Sobel [93], Prewitt [94] and Kirsch [90]. In all of them each pixel in the image is convolved with both

kernels. One kernel responds maximally to a generally vertical edge and the other to a horizontal edge. The maximum value of the two convolutions is taken as the output value for that pixel. The kernels for the Sobel edge operator are shown in Table 3.3, whereas their results are presented in Fig. 3.12 c).

Table 3.3 The Sobel convolution kernels.

-1	0	1	1	2	1
-2	0	2	0	0	0
-1	0	1	-1	-2	-1

The kernels for the Prewitt edge operator are shown in Table 3.4.

Table 3.4 Prewitt convolution kernels

-1	0	1	1	1	
-1	0	1	0	0	
-1	0	1	-1	-1	

The Canny edge detector [95] is known to many as the optimal one for a number of reasons; the first and most obvious being its low error rate. It is important that edges presented in images should not be missing and that there are no responses to non-edges. The second reason is that the edge points are well localized, while the third that there is only one response to a single edge.

In the Canny algorithm the Gaussian filter, based on the 5×5 mask, is used to smooth the image because the larger the Gaussian mask, the lower the detector's rate of sensitivity to noise. Next, the Sobel operator is applied to estimate the gradient Gx in the x-direction and Gy in the y-direction. The magnitude, the so-called *edge strength* of the gradient is thus approximated, using the formula:

$$|G| = |Gx| + |Gy|$$
(3.22)

Then we find the edge direction

$$\theta = \arctan\left(\frac{Gy}{Gx}\right) \tag{3.23}$$

See the result of the Canny edge operator applying to one segmented colour layer in Fig. 3.12 d).

We also use this algorithm to create contour map for the whole image as it is seen in Fig. 3.2.

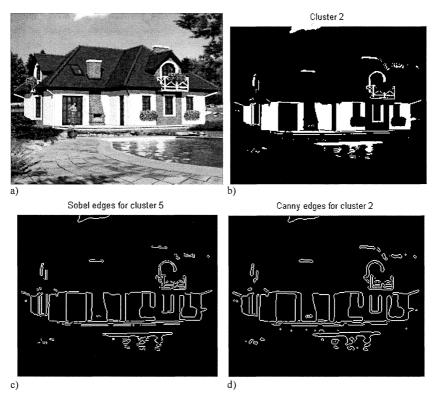


Fig. 3.12 An example of edge detection. a) the original image, b) a layer segmented by clustering, c) an example of the Sobel method for the layer from b), d) an example of the Canny method for the layer from b).

3.4.2 Boundary Tracking by Active Contours

In contrast to gradient-based representation, where the boundaries are detected based on the pixel intensity, the parametric model of active contours leads to the energy minimization problem.

Definition 3.2. (active contours)

Formally, let ρ be a metric (for instance the Euclidean metric) in \mathbf{R}^2 and $K(\mathbf{x}_0, \varepsilon) = \{\mathbf{x} \in \mathbf{R}^2 : \rho(\mathbf{x}_0, \mathbf{x}) < \varepsilon\}$ be a sphere with the centre $\mathbf{x}_0 \in \mathbf{R}^2$ and radius $\varepsilon < 0$. A set $c \subseteq \mathbf{R}^2$ is a contour if and only if there exists a function $f: \mathbf{R}^2 \to \mathbf{R}$, such as: $c = \{x \in \mathbf{R}^2 : \bigvee \land f(x_1) \ge 0 \land f(x_2) < 0\}$. $C(p) = \begin{bmatrix} x(p) \\ y(p) \end{bmatrix}$, where $p \in [0,1]$, that moves through the spatial domain Ω of an

image I(x,y). A snake, which we call the gradient vector flow (GVF) snake, begins with the calculation of a field of forces, called the GVF forces, over the image domain.

$$J(C) = E_{int}(C) + E_{ext}(C)$$
(3.24)

The external energy function E_{ext} is derived from the image so it moves towards the image contour:

$$E_{ext} = \int_{0}^{1} P(C(p)) \, dp = -\nabla P(C(p)) \tag{3.25}$$

where P(x,y) is a convolution of image I(x,y) (seen as a line) with a 2D Gaussian function $G_{\sigma}(x,y)$ with a standard deviation σ , as follows:

$$P(x,y) = -|\nabla G_{\sigma}(x,y) * I(x,y)|^{2}$$
(3.26)

The internal energy E_{int} controls the snake like a physical object resistant to both stretching and bending, towards the image boundaries:

$$E_{int} = \frac{1}{2} \int_{0}^{1} \alpha |C'(p)|^{2} + \beta |C''(p)|^{2} dp \qquad (3.27)$$

where the first derivative C'(p) models stretching and elasticity, whereas the second derivative C''(p) models bending and rigidity, where α and β are weight parameters.

A snake that minimizes J(C) must satisfy the Euler equation:

$$\alpha C''(p) - \beta C'''(p) - \nabla P(C(p)) = 0$$
(3.28)

that can be viewed as a force balance equation

$$F_{int} + F_{ext}^{(p)} = 0 \tag{3.29}$$

where: $F_{int} = \alpha C''(p) - \beta C'''(p)$ and $F_{ext}^{(p)} = -\nabla E_{ext}$.

The GVF forces create the gradient of an image edge map (see Fig. 3.13).

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Fig. 3.13 A gradient vector flow (GVF) field for a U-shaped object. These vectors will pull an active contour towards the object boundary. (Follows: *Active Contours, Deformable Models, and Gradient Vector Flow* Chenyang Xu and Jerry L. Prince web page: http://www.iacl.ece.jhu.edu/static/gvf/)

In comparison with the classical edge detection techniques, snakes have multiple advantages:

- They produce closed and smooth object boundaries.
- They autonomously and adaptively search for the minimum state.
- External image forces act upon the snake in an intuitive manner.
- Incorporating Gaussian smoothing in the image energy function introduces scale sensitivity.

But they also have some key drawbacks:

- They are sensitive to local minima states.
- Minute features are often ignored during energy minimization over the entire contour.
- Their accuracy depends on the convergence policy.

#### 3.4.3 Hough Transform

The classical Hough transform was concerned with the identification of lines in the image, but later the Hough transform was extended to identify the positions of arbitrary shapes, most commonly circles or ellipses.

The simplest variant of the Hough transform is used to detect of straight lines. In general, the straight line y = mx + b can be represented as a point (b,m) in the parameter space. However, vertical lines pose a problem. Instead, Duda and Hart [97] propose the polar coordinate representation of a line:

$$\rho = x \cos\theta + y \sin\theta, \tag{3.30}$$

where  $\rho$  is the distance from the origin to the closest point on the straight line, and  $\theta$  is the angle between the x axis and the line connecting the origin with that closest point (see Fig. 3.14).

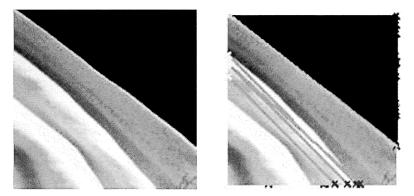


Fig. 3.14 Left: The original image. Right: Lines (green) found by the Hough transform.

Each point in image generates the sinusoid in Hough space (Fig. 3.15), and each point along this sinusoid corresponds to the  $\rho$ - $\theta$  values for a single line passing through the original point.

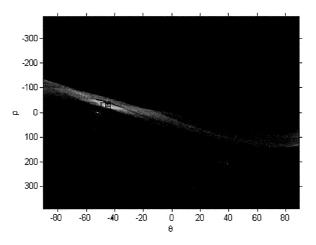


Fig. 3.15 The Hough transform space. White sinusoids represents lines visible in Fig. 3.14.

The Hough space is an accumulator space which means that it sums up the votes of many pixels in the image, and points in Hough space that have a large total vote are then interpreted as indicating the corresponding alignment on the real space image. To construct the Hough transform, every point present in the real-space image casts its votes into the Hough space for each of the lines that can possibly pass through it.

#### 3.5 Shape Information

Shape extraction is a non-trivial operation, but shape-based methods are particularly challenging due to the intrinsic difficulties in dealing with shape location and recognition. Nevertheless, there is no doubt that shape is one of the basic features describing image content, hence, we define the key properties of shape:

- identifiability: shapes which are found perceptually similar by humans have the same/analogous features that are different from the others;
- translation, rotation, scale and affine invariance;
- the location, the rotation and the scaling changes of the shape must not affect the extracted features;
- occultation invariance: when some parts of a shape are occulted by other objects, the feature of the remaining part must not change compared to the original shape.
- statistically independent of two features. This represents compactness of the representation.
- reliability: as long as one deals with the same pattern, the extracted features must remain the same.

Shape description can be generally divided into two kinds of methods: contourbased and region-based. Under each kind, the methods are further divided into a structural and global approach based on whether the shape is represented as a whole or by segments (primitives). The whole breakdown is shown in Fig. 3.16.

Contour shape techniques only exploit shape boundary information. Zhu et al. [98] use salient contours, extracted from bottom-up contour grouping, as tokens for image-model shape matching. Shape matching with contours instead of isolated edges has several advantages. Long salient contours are more distinctive, which leads to efficiency of the search as well as the accuracy of shape matching. Furthermore, an accidental alignment causing false positive detections is removed by requiring the entire contour to match whole objects.

#### Shape describing methods

Contour-based		Region-based	
structural chain code	global - perimeter	structural (	⊐ global  - area
• polygon	<ul> <li>compactness</li> </ul>	+ media axis	• Euler number
- B-spline	- eccentricity	• core	- eccentricty
→ invariants	→ shape signature	F	· geometric moments
	→ Hausdorff distance		· Zernike moments
	<ul> <li>Fourier descriptor</li> </ul>		
	• wavelet descriptor		Legendre moments
	→ curvature scale space		⁺ grid method
	+ autoregressive		+ shape matrix
	- elastic matching		1

Fig. 3.16 Shape describing methods [99].

Boundary-based methods such as [100] represent shapes by the locations of the maxima of its curvature scale space (CSS) image. Shapes are smoothed by selecting the appropriate scale and then matched by shifting the CSS contours so that the major maxima of one image overlap that of the other [101]. The shape boundaries are approximated using planar curves and are progressively simplified through discrete curve evolution based on a novel relevance measure [100]. The weakness of the boundary-based approach is that it does not represent the interior of the shape [102] and is, therefore, very sensitive to spatial reconfigurations of parts and local boundary perturbations.

In region-based techniques, all the pixels within a shape region are taken into account to obtain the shape representation. Common region based methods use moment descriptors to describe shapes [103]. Other region based methods include the grid method, shape matrix, convex hull and media axis.

## 3.5.1 The Shape Approach to the Hybrid Semantic System

According to the shape descriptors presented above our approach belongs to the region-based group of methods (see Fig. 3.16). Generally, we define shape descriptors for separated segments. The method used to find these segments is introduced in section 4.2.3.

Assuming that we have segments described as a set of pixels, we can apply two types of moments: moments of inertia and Zernike moments. The former are very efficient as shape descriptors and can be calculated as:

$$\mu_{pq} = \sum_{x} \sum_{y} (x - \bar{x})^p (y - \bar{y})^q I(x, y), \qquad p, q = 0, 1, 2$$
(3.31)

where p and q are the number of moments,  $\bar{x}$  and  $\bar{y}$  are segment centroids.

The second most efficient kind of descriptors is Zernike moments [104].

Definition 3.3 (Zernike moments)

Zernike moments are a set of complex polynomials  $\{V_{pq}(x,y)\}$  which form a complete orthogonal set over the unit disk of  $x^2 + y^2 \le 1$ . Hence, the definition of 2D Zernike moments with  $p^{\text{th}}$  order with repetition q for intensity image I(x,y) of the image is described as:

$$Z_{pq} = \frac{p+1}{\pi} \iint_{x^2 + y^2 \le 1} V_{pq}^*(x, y) I(x, y) dx dy$$
(3.32)

where:  $V_{pq}^{*}(x, y) = V_{p,-q}(x, y).$  (3.33)

Generally, the first 10 Zernike moments, i.e. those from  $Z_{00}$  to  $Z_{33}$ , are sufficient as a shape feature (see Fig. 3.17). The scale invariance is obtained by/thanks to normalization of  $Z_{00}$  by the total number of image pixels.

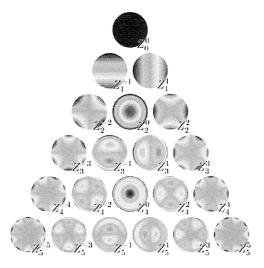


Fig. 3.17 The first Zernike base functions (followed Wikipedia).

Characteristic features of Zernike moments are: (i) invariance to rotation only; (ii) the translation invariance is achieved by the location of the original image centroid in the centre of the coordinates, (iii) the scale invariance is obtained by normalizing  $Z_{00}$  by the total number of image pixels.

Although the Zernike moment descriptor has a robust performance, it has several shortcomings. First, the kernel of Zernike moments is complex to compute, and the shape has to be normalized into a unit disk before deriving the moment features. Second, the radial features and circular features captured by Zernike moments are not consistent, one is in the spatial domain and the other is in the spectral domain. It does not allow multi-resolution analysis of a shape in a radial direction. Third, the circular spectral features are not captured evenly at each order, which can result in a loss of significant features which are useful for shape description.

To overcome these shortcomings, a generic Fourier descriptor (GFD) has been proposed by Zhang and Lu [99]. The GFD is acquired by applying a 2D Fourier transform on a polar-raster PF:

$$PF = \sum_{r} \sum_{i} I(r, \theta_i) \exp\left[2\pi j \left(\frac{r}{R}\rho + \frac{2\pi i}{T}\varphi\right)\right]$$
(3.34)

where:  $0 \le r < R$  and  $\theta_i = i(2\pi/T)$ ;  $0 \le i < T$ ;  $0 \le \rho < R$ ,  $0 \le \varphi < T$ . *R* and *T* are the radial frequency resolution and angular frequency resolution, respectively. The normalized coefficients are the GFD. The similarity between two shapes is measured by the city block distance between their GFDs.

It has been found that methods operating within the spatial domain suffer from two main drawbacks: noise sensitivity and a high dimension of the feature vector. The problems can be solved in four ways: histogram, moments, scale space and spectral transforms.

## **3.6 Local Feature Descriptors**

Local feature descriptors represent a group of methods which allow the user to find local image structures in a repeatable way and to encode them in a representation that is invariant to a range of image transformations, such as translation, rotation, scaling, and affine deformation. The purpose of introducing local feature descriptors is to provide a representation that enables the user to efficiently match local structures of images. For this objective, the feature extractors must fulfil three important criteria:

- 1. The feature extraction process should be repeatable and precise, so that the same features are extracted on two images showing the same object.
- 2. At the same time, the features should be distinctive, so that different image structures can be held apart from each other.

3. Proper feature descriptors should be resistant to accidental variance of features and invariant to scaling and rotation of image.

Herein, we present a group of seven local feature descriptors developed recently. Their general advantage is high precision in matching the required object, but their drawback is long running time and the necessity to provide a query-by-example.

## 3.6.1 Scale-Invariant Feature Transform (SIFT)

The scale-invariant feature transform (SIFT) was introduced by Lowe [24], [23] to identify objects in two images, even if these objects were cluttered or under partial occlusion, because the SIFT feature descriptor is invariant to uniform scaling and orientation. In like manner, it is partially invariant to affine distortion and illumination changes.

The algorithm starts from key-points detection in order to identify locations and scales that can be repeatedly assigned under differing views of the same object. Key-point locations are defined as maxima and minima of the difference of the Gaussians  $G(x, y, \sigma)$  applied in a scale-space to a series of smoothed and resampled images.

$$L(x, y, \sigma) = G(x, y, \sigma) * I(x, y)$$
(3.35)

where  $L(x,y,\sigma)$  is the product of a convolution. This difference of Gaussians is calculated for two nearby scales with k factor:

$$D(x, y, \sigma) = L(x, y, k\sigma) - L(x, y, \sigma)$$
(3.36)

The scale-space extrema detection produces too many key-point candidates, so at first, for each candidate key-point, interpolation of nearby data is used to accurately determine its position. The interpolation is done using the quadratic Taylor expansion of the difference-of-Gaussian scale-space function, D with the candidate key-point as the origin. This Taylor expansion is given by:

$$D(\mathbf{x}) = D + \frac{\partial D^{T}}{\partial \mathbf{x}} \mathbf{x} + \frac{1}{2} \mathbf{x}^{T} \frac{\partial^{2} D}{\partial \mathbf{x}^{2}} \mathbf{x}$$
(3.37)

for  $\mathbf{x} = (x, y, \sigma)^{\mathrm{T}}$ .

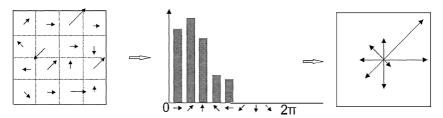
Low contrast candidate points and edge response points along an edge are discarded. Dominant orientations are assigned to localized key-points. These steps ensure that the key-points are more stable for matching and recognition. The SIFT descriptors robust to local affine distortion are then obtained by considering pixels around a radius of the key location, blurring and resampling of local image orientation planes.

The next step is the orientation assignment when each key-point is assigned one or more orientations based on local image gradient directions. First, the Gaussian-smoothed image  $L(x,y,\sigma)$  at the key-point's scale  $\sigma$  is taken so that all computations are performed in a scale-invariant manner. For an image sample L(x,y) at scale  $\sigma$ , the gradient magnitude m(x,y) and orientation  $\theta(x,y)$  are precomputed using pixel differences:

$$m(x,y) = \sqrt{(L(x+1,y) - L(x-1,y))^2 + (L(x,y+1) - L(x,y-1))^2} \quad (3.38)$$

$$\theta(x,y) = \arctan\left(\frac{L(x,y+1) - L(x,y-1)}{L(x+1,y) - L(x-1,y)}\right)$$
(3.39)

assuming that  $\forall (x, y)$  in the neighbourhood  $(x_0, y_0)$ .



**Fig. 3.18** The gradient magnitude and orientation at each point of a 4x4 set of samples (on the left) which are accumulated into orientation histograms with 8 bins each (in the middle). The key-points descriptor summarizes the contents over 4x4 subregions, as shown on the right, with the length of each arrow corresponding to the sum of the gradient magnitudes near that direction within the region. Peaks in the orientation histogram correspond to dominant directions of local gradients.

The SIFT key samples generated at a larger scale are given twice the weight of those at a smaller scale. This means that the larger scale is in effect able to filter the most likely neighbours for the smaller scale. This also improves recognition performance by giving more weight to the least-noisy scale. To avoid the problem of boundary effects in bin assignment, each key-point match votes for the 2 closest bins in each dimension, giving a total of 16 entries for each hypothesis and further broadening the pose range.

Hough transform (as it has been described in subsect. 3.4.3, cf. (3.30)) is used to cluster reliable model hypotheses to search for keys that agree upon a particular model pose. When feature clusters are found to vote for the same pose of an object, the probability of the interpretation being correct is much higher than for any single feature. An entry in a hash table is created predicting the model location, orientation, and scale from the match hypothesis. The hash table is searched to identify all clusters of at least 3 entries in a bin, and the bins are sorted into decreasing order of size.

Each identified cluster is then subject to a verification procedure in which a linear least squares solution is performed for the parameters of the affine

transformation relating the model to the image. The affine transformation of a model point  $[x y]^{T}$  to an image point  $[u v]^{T}$  can be written as below:

$$\begin{bmatrix} u \\ v \end{bmatrix} = \begin{bmatrix} m_1 & m_2 \\ m_3 & m_4 \end{bmatrix} \begin{bmatrix} x \\ y \end{bmatrix} + \begin{bmatrix} t_x \\ t_y \end{bmatrix}$$
(3.40)

where the model translation is  $[t_x t_y]^T$  and the affine rotation, scale, and stretch are represented by the parameters  $m_1$ ,  $m_2$ ,  $m_3$  and  $m_4$ . In order to find the transformation parameters the equation (3.40) can be reformulated to group the unknowns into a column vector.

$$\begin{bmatrix} x & y & 0 & 0 & 1 & 0 \\ 0 & 0 & x & y & 0 & 1 \\ & \dots & & \\ & & \dots & & \\ & & & \dots & & \end{bmatrix} \begin{bmatrix} m_1 \\ m_2 \\ m_3 \\ m_4 \\ t_x \\ t_y \end{bmatrix} = \begin{bmatrix} u \\ v \\ \vdots \end{bmatrix}$$
(3.41)

This equation presents a single match, but any number of further matches can be added, with each match contributing two more rows to the first and last matrix. At least 3 matches are needed to provide a solution. We can write this linear system as

$$A\mathbf{x} \approx \mathbf{b}$$
 (3.42)

where A is a known *n*-by-*p* matrix (n > p), **x** is an unknown *p*-dimensional parameter vector, and **b** is a known *n*-dimensional measurement vector. Therefore, the minimizing vector **x** is a solution of the normal equation.

$$\mathbf{A}^{\mathrm{T}}\mathbf{A}\mathbf{x} = \mathbf{A}^{\mathrm{T}}\mathbf{b} \tag{3.43}$$

hence, we obtain:

$$\mathbf{x} = (A^T A)^{-1} A^T \mathbf{b} \tag{3.44}$$

which minimizes the sum of the squares of the distances from the projected model locations to the corresponding image locations.

# 3.6.2 RootSIFT

SIFT was originally designed, by Lowe [23], to be used with the Euclidean distance, but since there is a histogram comparison, Arandjelović and Zisserman

[105] introduced alternative histogram distance measures, namely the Hellinger kernel.

The Hellinger kernel for two L₁ normalized histograms, x and y (i.e.  $\sum_{i=1}^{n} x_i = 1$  and  $x_i \ge 0$ ), is defined as follows:

$$H(x,y) = \sum_{i=1}^{n} \sqrt{x_i y_i}$$
(3.45)

where *n* is a number of vector with unit Euclidean norm such as:  $\|\mathbf{x}\|_2 = 1$ .

The RootSIFT application slightly increases the average precision of retrieval.

#### 3.6.3 Rotation-Invariant Generalization of SIFT (RIFT)

RIFT is a rotation-invariant variant of the SIFT method dedicated to texture images where the notion of orientation is difficult to define. The RIFT descriptor is constructed using circular normalized patches divided into concentric rings of equal width and within each ring a gradient orientation histogram is computed [106], [55]. To maintain rotation invariance, the orientation is measured at each point relative to the direction pointing outward from the center (see Fig. 3.19).

When the size of the Laplacian-of-Gaussian (LoG) kernel matches with the size of a blob-like structure, the response attains an extremum:

$$|\text{LoG}(\mathbf{x}, \sigma_n)| = \sigma_n^2 |L_{xx}(\mathbf{x}, \sigma_n) + L_{yy}(\mathbf{x}, \sigma_n)|.$$
(3.46)

The LoG kernel can therefore be interpreted as a matching filter.

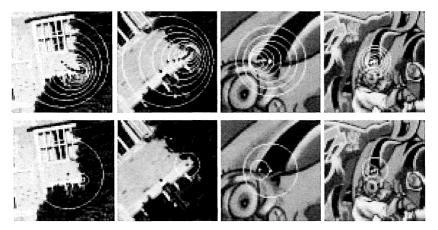


Fig. 3.19 Scale invariant interest point detection: (Top) Initial multi-scale Harris points (selected manually) corresponding to one local structure [106].

#### 3.6.4 Fisher Vector (FV)

The Fisher kernel has been proposed, in the context of measuring the amount of information that an observable random variable X carries about an unknown parameter  $\lambda$  of a distribution that models X. Formally, it is the variance of the score, or the expected value of the observed information. From the theory of information we know that the entropy of a random value V is:

$$H(V) = -\sum p(v) \log(p(v)) \tag{3.47}$$

In the context of image retrieval the FV are usually  $\ell^2$ -normalized since, as proved it Perronnin et al. [107] this is a way to eliminate the fact that distinct images contain different amounts of image-specific information. The Fisher Vector is applied to non-binary local features, using the Gaussian Mixture Model to represent the average distribution  $p(X | \lambda)$ .

For sample  $X = \{x_t, t = 1,...,T\}$  of observations, which is a set of *T* local descriptors extracted from an image and  $p(X | \lambda)$  represents a probability density function with  $\lambda$  parameters, the gradient  $\nabla_{\lambda}$  of the log-likelihood describes the contribution of the parameters to the generation process [107]:

$$G(X|\lambda) = \frac{1}{T} \nabla_{\lambda} \log p(X|\lambda)$$
(3.48)

The dimensionality of this vector depends only on the number of parameters  $\lambda$ , not on the number of patches T.

Further, this approach has been moved to an image classification, where the Fisher kernel K method is to derive a function that measures the similarity between two sets of data X and Y, such as the sets of local descriptors extracted from two images. The idea is to characterize a signal with a gradient vector derived from a probability density function (pdf) which models the generation process of the signal [108].

$$K(X,Y) = \mathbf{G}'(X|\lambda) \,\mathbf{F}^{-1}(\lambda) \,\mathbf{G}(Y|\lambda) \tag{3.49}$$

where Fisher information matrix:  $F^{-1}(\lambda) = L'(\lambda)L(\lambda)$ .

This representation can then be used as input to a discriminative classier. For the problem of image categorization the input signals are images. Perronnin and Dance proposed to use, as a generative model the Gaussian Mixture Models (GMM) which approximates the distribution of low-level features in images, i.e. a visual vocabulary.

Krapac and Segvić proposed in [109] to use large FVs to object location on video images. They detect multiple objects on complex backgrounds, for instance, road signs.

## 3.6.5 Vectors of Locally Aggregated Descriptors (VLAD)

The VLAD method analyzes the local descriptors contained in an image to create statistical summaries that still preserve the effectiveness of local descriptors and allow treating them as global descriptors [110]. These image descriptors were designed to be very low dimensional (e.g. 16 bytes per image).

This method encodes a set of local feature descriptors  $F = (x_1, ..., x_k)$ , extracted from an image treated as a codebook with k visual words, using a dictionary based on a clustering method, such as GMM or k-means clustering. Each local descriptor  $x_i$  is then associated with its nearest centroid NN $(x_i) = \mu_i$ .

$$v_{i,j} = \sum_{x_j: NN(x_j) = \mu_i} (x_j - \mu_{i,j})$$
(3.50)

where i=1,...,k is the index of visual word (k – number of centroids) and j=1,...,d is the local descriptor component. Hence, the whole image representation dimension is  $D = k \times d$ .

For each cluster, the residual vectors (i.e. the difference between the centroid and the associated descriptors) are accumulated and the sum of the residual is concatenated into a single vector  $V = [v_1^T \dots v_k^T]$ . Next, vector v is normalised by  $v := v/||v||_2$  and the Euclidean distance is sufficient to compare two VLADs.

#### 3.6.6. Features from accelerated segment test (FAST)

FAST is a corner detection method, introduced by Rosten and Drummond in 2006 [111], which could be used to extract feature points and later used to track and map objects [112]. A FAST corner detector uses a circle of 16 pixels (a Bresenham circle of r=3) to classify whether a candidate point p is actually a corner. Each pixel in the circle is labelled from integer number 1 to 16 clockwise. For a set of N contiguous pixels, if the pixels in the circle are all brighter than the intensity of candidate pixel p (denoted by  $I_p$ ), plus a threshold value t, or all darker than the intensity of candidate pixel p, minus threshold value t, then p is classified as a corner. The conditions can be written as:

- 1. A set of N contiguous pixels S,  $\forall x \in S$ , the intensity of x denoted by  $(I_x)$  can be  $I_x > I_p + t$ ;
- 2. A set of N contiguous pixels S,  $\forall x \in S, I_x \leq I_p t$ ;

So when either of the two conditions is met, candidate p can be classified as a corner. There is a tradeoff between selecting N, the number of contiguous pixels and the threshold value t. Then, N is usually selected as 12. A high-speed test method could be applied to exclude non-corner points.

Generally, the FAST detector is employed to find objects in video frames because of its effectiveness.

#### 3.6.7 Oriented FAST and Rotated BRIEF (ORB)

ORB is basically a fusion of the FAST key-point detector and a BRIEF descriptor with many modifications to enhance the performance introduced by Rublee et al. [113] in 2011. First, it uses FAST to find key-points, then apply the Harris [114] corner measure to find top N points among them. It also uses a pyramid to produce multiscale-features.

In order to compute orientation, they found moments of order p and q, such as:

$$m_{pq} = \sum_{x} \sum_{y} x^{p} y^{q} I(x, y), \quad p, q = 0, 1, 2$$
(3.51)

and intensity centroids:  $C = \left(\frac{m_{10} \ m_{01}}{m_{00} \ m_{00}}\right)$ .

Then, a vector  $\overrightarrow{OC}$  from the corner's center, O, to the centroid, can be constructed. The orientation of the patch then simply is:

$$\theta = \operatorname{atan2}(m_{01}, m_{10}), \tag{3.52}$$

where atan2 is the arctangent function with two arguments. In order to improve the rotation invariance of this measure authors made sure that moments are computed with x and y remaining within a circular region of radius r. They empirically selected r to be the patch size, so that that x and y run from [-r, r].

Next, the Binary Robust Independent Elementary Features (BRIEF) descriptor is used for a simple binary test  $\tau$  between pixels in a smoothed image patch **p**, as follows:

$$\tau(\mathbf{p}; \mathbf{x}, \mathbf{y}) \coloneqq \begin{cases} 1: \ \mathbf{p}(\mathbf{x}) < \mathbf{p}(\mathbf{y}) \\ 0: \ \mathbf{p}(\mathbf{x}) \ge \mathbf{p}(\mathbf{y}) \end{cases}$$
(3.53)

where  $\mathbf{p}(x)$  is the intensity of  $\mathbf{p}$  at a point *x*. The feature is defined as a vector of *n* binary tests:

$$f_n(\mathbf{p}) := \sum_{1 \le i \le n} 2^{i-1} \tau(\mathbf{p}; \mathbf{x}_i, \mathbf{y}_i)$$
(3.54)

The test pairs of x and y are selected by Gaussian distribution around the centre of the patch or PCA for good discrimination. As tests for typical frames of size  $640 \times 480$  proved, the ORB descriptor gives significant time decreasing.

## 3.7 Standardization Efforts - MPEG-7

The Moving Picture Experts Group (MPEG) [115] is a working group of authorities that was formed by ISO and IEC to set standards for audio and video compression and transmission. The MPEG-7 is a multimedia content description interface that uses XML to store metadata, and can be attached to the timecode in order to tag particular events. The description of the standard can be found in [116].

The main elements of the MPEG-7 standard are:

- Descriptors (D) that define the syntax and the semantics of each feature (metadata element);
- Description Schemes (DS) that specify the structure and semantics of the relationships between their components which may be both descriptors and description schemes;
- A Description Definition Language (DDL) to define the syntax of the MPEG-7 Description Tools and to allow the creation of new description schemes and, possibly, descriptors and to allow the extension and modification of existing description schemes;

The above-mentioned tools deal with binarization, synchronization, transport and storage of descriptors (see Fig. 3.20) [117].

MPEG-7 Visual Description Tools consist of basic structures and descriptors that cover the following basic visual features: color, texture, shape, motion, localization, and face recognition. Each category consists of elementary and sophisticated descriptors.

MPEG-7 Multimedia Description Schemes (DSs) are metadata structures for describing and annotating audio-visual (AV) content. The DSs provide a standardized way of describing in XML the important concepts related to AV content description and content management in order to facilitate searching, indexing, filtering, and access. The DSs are defined using the MPEG-7 Description Definition Language (DDL), which is based on the XML Schema language, and are instantiated as documents or streams. The resulting descriptions can be expressed in a textual form (i.e., human readable XML for editing, searching, filtering) or compressed binary form (i.e., for storage or transmission).

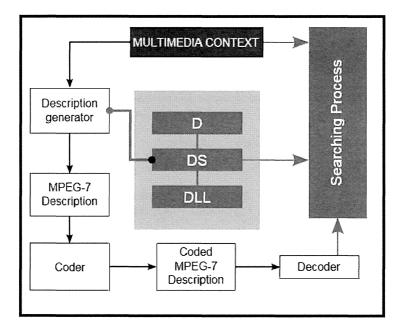


Fig. 3.20 Relations between different tools and the elaboration process of MPEG 7 [117].

## 3.8 Global Versus Local Comparison of Features

A feature is defined to capture a certain visual property of an image, either globally for the entire image or locally for a small group of pixels. So far, we have described local features i,e. related to the most basic structure of images, such as pixels. To reduce computation, an image may be divided into small, non-overlapping structures like lines, pixel neighbourhoods, subsets of pixels or patterns, and later the features are computed individually for every structure. The features are still local because of the small block size, but the amount of computation is radically reduced in comparison with that needed for obtaining features around every pixel.

In turn, global features capture the overall characteristics of an image. The advantage of global extraction is its high speed for both extracting features and computing similarity.

Both global and local features can be represented as a feature vector. There is one global feature for an image as well as many feature vectors  $\mathbf{x}$ ,  $\mathbf{y}$  with local features, also for this image, additionally, to each feature its weight  $\xi(i)$  can be

attributed. Then the Euclidean distance, measuring the distance between x and y vector, can be calculated as:

$$d_E = \sqrt{(\mathbf{x} - \mathbf{y})^T \operatorname{diag}(\xi_i^2)(\mathbf{x} - \mathbf{y})}$$
(3.55)

Generally, a histogram was widely used as a global feature because it is very fast to compute and easy to compare with another histogram. A histogram can then be treated as a k-dimensional vector  $(f_1, f_2, ..., f_k)$ , where  $f_i$  is the frequency of the  $l^{\text{th}}$  bin. We assume that two distributions (two images) have two histograms  $H_0(i)$  and  $H_1(i)$  with  $0 \le i \le n$  bins each.

We can, for example, compare histograms based on the Minkowski distance (with p=3):

$$d_{L_p}(H_0, H_1) = \left[\sum_{i=1}^n |H_0(i) - H_1(i)|^p\right]^{\frac{1}{p}}$$
(3.56)

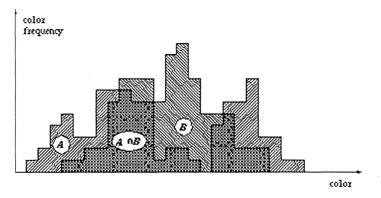


Fig. 3.21 Histogram intersection.

This kind of distance is very sensitive to even small shifts because the same bars must be compared with each other.

The histogram intersection [118] is widely used because of its ability to handle partial matches when the areas of the two histograms (the sum over all the bins (3.57)) are different (see Fig. 3.21). It is shown by Swain and Ballard [118] that when the areas of the two histograms are equal, the histogram intersection is equivalent to the (normalized)  $L_1$  distance.

$$d_{\cap}(H_0, H_1) = 1 - \frac{\sum_i \min(H_0(i) - H_1(i))}{\sum_i H_1(i)}$$
(3.57)

When the feature vector represents relative frequency distribution (e.g., a normalised grey level co-occurrence histogram), for example, texture features, the

dissimilarity can also be measured by the relative entropy, or Kullback-Leibler (K-L) divergence. Let  $D_J$  denote the divergence between two distributions,  $H_0(i)$  and  $H_1(i)$ , which is based on vector quantization. Then:

$$D_J(H_0, H_1) = \sum_{i=1}^n H_0(i) \log \frac{H_0(i)}{H_1(i)}$$
(3.58)

This dissimilarity measure is asymmetric and does not represent a distance because the triangle inequality is not satisfied. The symmetric distance is obtained by averaging  $H_0$  and  $H_1$  [119]. In histogram notation we can describe this distance as:

$$D_J(H_0, H_1) = \sum_{i=1}^n \left[ H_0(i) \log \frac{H_0(i)}{m_i} + H_1(i) \log \frac{H_1(i)}{m_i} \right]$$
(3.59)

where:  $m_i = \frac{H_0(i) + H_1(i)}{2}$ . Additionally, the K-L divergence is sensitive to histogram binning.

In 1998 Rubner et al. [12] introduced the Earth Mover's Distance (EMD) which was understood as the minimal cost that must be paid to transform one distribution (histogram) into the other, where there is a *distance*  $d_{i,j}$  (between bin *i* and *j*) and is meant as the distance between the basic features that are aggregated into the histogram. Computing the EMD is based on a solution to the well-known *transportation problem*.

Given two histograms  $H_i$  and  $H_j$ , the EMD is:

$$EMD(H_{i}, H_{j}) = \frac{\min_{f_{i,j}} \sum_{i,j} (f_{i,j} d_{i,j})}{\sum_{i,j} (f_{i,j})}$$
(3.60)

for the following constraints:  $f_{i,j} \ge 0$ ,  $\sum_j f_{i,j} \le H_i$ ;  $\sum_i f_{i,j} \le H_j$ ;

$$\sum_{i,j} (f_{i,j}) = \min \left\{ \sum_i (H_i) , \sum_j (H_j) \right\}$$

where:  $f_{i,j}$  denotes the flow which allows us to move some amount of 'mass' from  $H_i$  to  $H_j$  and vice versa; the flow cannot be higher than neither  $H_i$  nor  $H_j$  and the last constraint describes the maximum mass possible to move.

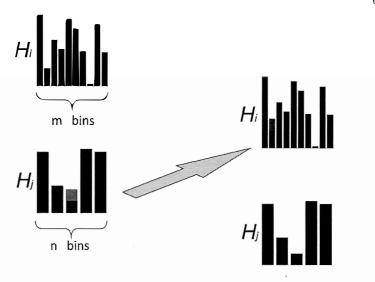


Fig. 3.22 Transport of 'mass' from H_i to H_i.

The EMD naturally extends the notion of a distance between individual elements to that of a distance between sets, or distributions, of elements. The advantages of the EMD over previous definitions of distribution distances are significant.

If the ground distance is a metric and the total weights of two signatures are equal, the EMD is a true metric, which allows endowing image spaces with a metric structure.

#### **3.8** From Features to Signature

Generally, a signature describes the image information (also known as a global image descriptor) and it can also be seen as a mathematical function. The principal purpose of this function is to extract from a large image data structure.

The objective is that computed signatures enable us to determine similarities (i.e have matching features) between images that represent, for instance, the same scene but captured from different points of view. It means that most of the visual applications required that two images with high perceptual similarities have resembling signatures.

Signature generation functions can be roughly classified in three main groups, depending on the input data used to generate the global descriptor:

1. Appearance-based: the signature is calculated from texture, colour information, transformations in the frequency space or matrix factorizations .

- 2. Feature-based: the signature is calculated from the image key-points and their descriptors like those used in the SIFT, RIFT, etc.
- 3. Region-based: the signature is defined on the distance between sets of vectors, which is not as obvious as defining distance between single vectors.

A signature  $\{\mathbf{s}_j = (\mathbf{m} \ j; \ w\mathbf{m} \ j)\}$  can represent a set of feature clusters, where a cluster is represented by its mean (or mode)  $m_j$ , and by the fraction  $w\mathbf{m} \ j$  of pixels that belong to that cluster,  $1 \le j \le n$ , where *n* depicts the complexity of the particular image and the representative  $m_j$  is a *d*-dimensional vector. In general, vector quantization algorithms can be used for the clustering, as long as they are applied on every image independently, and they adjust the number of clusters to the complexities of individual images. On this assumption, a histogram  $\{h_j\}$  can be viewed as a signature  $\{\mathbf{s}_j = (\mathbf{m} \ j; \ w\mathbf{m} \ j)\}$  in which  $j^{\text{th}}$  cluster maps the point  $m_j$  with the central value in bin j of the histogram, and then  $w_j$  is equal to  $h_j$ , which better represents the image content.

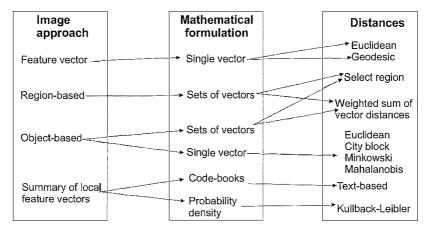


Fig. 3.23 Different types of image similarity measure and their mathematical formulations.

Additionally, the global signature based on the image histogram can be seen as:  $\{(z'_1, f'_1), ..., (z'_k, f'_k)\}$ , where  $f'_l$  is the percentage of  $x_{i,j}$ 's grouped into cluster l, and  $z'_l$  is a bin's location. The collection of pixels (i, j) for which  $x_{i,j}$ 's are in the same cluster forms a relatively homogeneous region because the common cluster forces closeness between the visual features in  $x_{i,j}$ 's.

As shown in Fig. 3.23, similarity computation can be performed with feature vectors, region-based signature, object-based signature, or summarized local features.

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