



# Graphs for image processing, analysis and pattern recognition

Florence Tupin

### Overview

#### 1. Definitions and representation models

#### 2. Single graph methods

- Segmentation or labeling and graph-cuts
- Graphs for pattern recognition

### 3. Graph matching

- Graph or subgraph isomorphisms
- Error tolerant graph-matching
- Approximate algorithms (*inexact matching*)

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# Why using graphs ?

- Interest: they give a compact, structured and complete representation, easy to handle
- Applications:

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- Image processing: segmentation, boundary detection
- Pattern recognition: printed characters, objects (buildings 2D ou 3D, brain structures, ...), faces, ...

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- Image registration
- Understanding of structured scenes

### Definitions

Graph: 
$$G = (X, E)$$

- X set of nodes (|X| order of the graph)
- E set of edges (|E| size of the graph)
- complete graph (size  $\frac{n(n-1)}{2}$ )
- partial graph G = (X, E') with E' part of E
- ▶ subgraph F = (Y, E'),  $Y \subseteq X$  et  $E' \subseteq E$
- degree of a node x : d(x) = number of edges
- connected graph: for each pair of nodes you find a path linking them
- tree: connected graph without cycle
- clique: complete subgraph
- dual graph (face  $\rightarrow$  node)
- segment graph (edge  $\rightarrow$  node)
- hypergraph (n-ary relations)
- weighted graphs: weights on the edges



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with  $D_{ii} = d_i$  (D degree matrix)

### Representation

Adjacency matrix, adjacency lists





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### Representation

Adjacency matrix, adjacency lists



#### FIGURE 1.4

From top-left to bottom-right: a weighted directed graph, its adjacency list, its adjacency matrix, and its (transposed) incidence matrix representations.

(figure from "Image processing and analysis with graphs", Lezoray - Grady)

Which graphs for images ? • X = enisbl. des pixels 1 E = 4- Connou B- CONN Att ributs des moeude -> ng, RVB, (Attributes des ares - (wij = Ingi-ngi) La radionnétriques ( Créj = ( Pi - Pi)) colorismétriques ( distances, relations apatiales...) ) glormétriques ( distances, relations apatiales...) • Graphes des primitives; Craphes des primitives; Craphes des régions : K-means K-means No Dr. > noeuds ~ régions , RAG · moyenne des ng/(downs arcs -> régions adjouentes / (GAR) • Mi Et postion du la postion

### Examples of graphs



### Examples of image graphs



#### FIGURE 1.11

The rectangular (left) and hexagonal (right) lattices and their associated Voronoi cells.





(c) 26-adjacency.

(a) 6-adjacency.

(b) 18-adjacency.

### Examples of image graphs



RAG (Region Adjacency Graph)

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### Examples of image graphs





(figure from "Image processing and analysis with graphs", Lezoray - Grady)

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### Examples of graphs

Graph of fuzzy attributes : attributed graph with fuzzy value for each attribute

Hierarchical graph :

multi-level graph and and bi-partite graph between 2 levels (multi-level approaches, object grouping, ...)

Exemples :

- quadtrees, octrees
- hierarchical representation of the brain

#### Graph for reasoning

decision tree, matching graph

### Graph example:



#### FIGURE 1.13

(a) An image with the quadtree tessellation, (b) the associated partition tree, (c) a real image with the quadtree tessellation, (d) the region adjacency graph associated to the quadtree partition, (e) and (f) two different irregular tessellations of an image using image-dependent superpixel segmentation methods: Watershed [23] and SLIC superpixels [24].

(figure from "Image processing and analysis with graphs", Lezoray - Grady)

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### Graph examples



**Figure 2** – Représentation de variété des points clés de  $S_{\omega}^{\max}(I)$  (en rouge) et  $S_{\omega}^{\min}(I)$  (en bleu image Pléiades ayant des textures locales différentes.

(figure from M.T. Pham PhD)

# Gra





(c) Détecteur de Harris



(b) Extrema locaux



### Graph examples



Figure 5 - Vecteur de description proposé pour l'analyse ponctuelle de la texture.

#### (figure from M.T. Pham PhD) (B) (E) (E) (E) (E) (C)

### Graph examples - BPT Binary Partition Tree



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Which algorithms from graph theory ?

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Mim Cut / Nax Hour

Min Spanning Tree

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Some classical algorithms

Search of the minimum spanning tree

• Kruskal algorithm  $O(n^2 + m \log_2(m))$ 

Prim algorithm O(n<sup>2</sup>)

Shortest path problems

• positive weights: Dijkstra algorithm  $O(n^2)$ 

• arbitrary weights but without cycle: Bellman algorithm  $O(n^2)$ Max flow and Min cut

► *G* = (*X*, *E*)

▶ partitioning in two sets A et B ( $A \cup B = X$ ,  $A \cap B = \emptyset$ )

• 
$$cut(A,B) = \sum_{x \in A, y \in B} w(x,y)$$

Ford and Fulkerson algorithm

Search of maximal clique in a graph

decision tree

cut of already explored branches

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Segmentation by minimum spanning tree  $\mathcal{W}_{\bar{\mathcal{U}}} = \langle \sigma g_i - \gamma g_i \rangle$ 

How can we segment this image using a minimum spanning tree ?

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Segmentation by minimum spanning tree

Constantinides (1986)

- graph of pixels weighted by the gray levels (or colors) (weights = distances)
- search of the minimum spanning tree
- ► spanning tree ⇒ partitioning by suppressing the most costly edges









image

graphe des pixels attribué

arbre couvrant de poids minimal

suppression des arêtes les plus coûteuses

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### Computation of the minimum spanning tree

### Kruskal algorithm

► Starting from a partial graph without any edge, iterate (n - 1) times : choose the edge of minimum weight creating no cycle in the graph with the previsouly chosen edges

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- In practice:
  - 1. sorting of edges by increasing weights
  - 2. while the number of edges is less than (n-1) do:
    - select the first edge not already examined
    - if cycle, reject
    - else, add the edge in the graph
- Complexity:  $O(n^2 + mlog_2(m))$

### Prim algorithm

- Extension from near to near of the current tree
- Complexity:  $O(n^2)$

# Constantinides (1986)









### Segmentation by graph-cut

Graph-cut definition:

- graph G = (X, E)
- ▶ partitioning in 2 parts A et B  $(A \cup B = X, A \cap B = \emptyset)$

• 
$$cut(A,B) = \sum_{i \in A, j \in B} w_{ij}$$



### Segmentation by graph clustering

Clustering : partitioning of the graph in groups of nodes based on their similarities [Each cluster : a closely connected component]

The clustering corresponds to:

- edges between different groups have low weights (weak similarities)
- edges inside a group have high weights (high similarities)

Possible cost functions for the cut:

- minimum cut  $Cut(A_1, ..., A_k) = \sum_{i=1}^{i=k} Cut(A_i, \overline{A_i})$
- minimum cut normalized by the size of each part (RatioCut) RatioCut(A<sub>1</sub>,...,A<sub>k</sub>) = ∑<sub>i=1</sub><sup>i=k</sup> 1/|A<sub>i</sub>| Cut(A<sub>i</sub>, A<sub>i</sub>) (|A<sub>i</sub>| number of vertices in A<sub>i</sub>)
   minimum cut normalized by the connectivity of each part (NCut) NCut(A<sub>1</sub>,...,A<sub>k</sub>) = ∑<sub>i=1</sub><sup>i=k</sup> 1/vol(A<sub>i</sub>) Cut(A<sub>i</sub>, A<sub>i</sub>)
  - $(vol(A_i) = \sum_{k \in A_i} d_k$  sum of the weight of all edges of

### Toy example



Influence of the number of edges: Cut(A, B) = 4b, Cut(A', B') = 3b



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### Normalized cut

- Principle: graph clustering
- + suppression of the influence of the number of edges: normalized cut

$$Ncut(A, B) = \underbrace{cut(A, B)}_{assoc(A, X)} + \underbrace{cut(A, B)}_{assoc(B, X)}$$
$$assoc(A, X) = \sum_{a \in A, x \in X} w(a, x) = vol(A)$$

Measuring the connectivity of a cluster:

$$Nassoc(A, B) = \frac{assoc(A, A)}{assoc(A, X)} + \frac{assoc(B, B)}{assoc(B, X)}$$
$$Ncut(A, B) = 2 - Nassoc(A, B)$$

minimizing the cut  $\Leftrightarrow$  maximizing group connectivity

### Toy example

Influence of the number of edges: Cut(A, B) = 4b, Cut(A', B') = 3b



 $\Rightarrow$  normalized cut (NCut)

vol(A) = 3a + 3a + 2b + 3a + 3a + 2b = 12a + 1b vol(B) = 3b + 3b = 6b.  $NCut(A, B) = \frac{4b}{12a + 4b} + \frac{4b}{6b} = \frac{4b}{12a + 4b} + \frac{2}{3}$  vol(A') = 12a + 4b + 3b = 12a + 7b vol(B') = 3b $NCut(A', B') = \frac{3b}{12a + 7b} + \frac{3b}{3b} = \frac{3b}{12a + 7b} + \frac{1}{12}$ 

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### Graph theory and cuts

#### MinCut by combinatorial optimization

- Stoer-Wagner algorithm
- Principle: iterative reducing of the graph by fusion of the nodes linked by the maximal weights

#### Min K-cut by combinatorial optimization

- Partitioning the (un-oriented graph) graph in many components
- Gomory-Hu algorithm

#### minCut in oriented graph by combinatorial optimization

- Ford-Fulkerson algorithm (oriented graph with two terminal nodes (sink / tank)
- Principle: MaxFlow search (MinCut equivalence) by search for an augmenting chain to increase the flow

### Graph theory and cuts

Laplacian matrices  $D = diag(d_i)$  with  $d_i = \sum_j w_{ij}$   $W = (w_{ij})$ Graph Laplacian matrix L = D - WNormalized graph Laplacian matrix  $L_n = D^{-\frac{1}{2}}LD^{-\frac{1}{2}} = I - D^{-\frac{1}{2}}WD^{-\frac{1}{2}}$ 

clustering en

#### Spectral clustering) algorithms and cuts

- Computation of the eigen-values and eigen-vectors of some matrix (L, L<sub>n</sub>, or generalized eigen problems Lu = λDu)
   selection of the k smallest eigen-values and associated k eigen-vectors u<sub>k</sub>
   U = (u<sub>1</sub>,..., u<sub>k</sub>) ∈ R<sup>n×k</sup>
   If y<sub>i</sub> ∈ R<sup>k</sup> be the ith row of U (i = 1, ..., n)
   cluster the points (y<sub>i</sub>) = w with the k means algorithm into
- ► cluster the points (y<sub>i</sub>)<sub>1≤i≤n</sub> with the k-means algorithm into clusters C<sub>1</sub>,..., C<sub>k</sub>
- clusters  $A_1, ..., A_k$  with  $A_i = \{j | y_i \in C_i\}$

Examples (univ. Berkeley) { noeud i (pixeli) + Pi, Ci noeud i (pixeli) + Pi, Ci



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http://www.cs.berkeley.edu/projects/vision/Grouping/

### Examples (univ. Berkeley)



http://www.cs.berkeley.edu/projects/vision/Grouping/
### Examples (univ. Berkeley)



http://www.cs.berkeley.edu/projects/vision/Grouping/ => <=> = > > >

# Examples (univ. Alberta) with linear constraints



### Examples (Mean Shift et Normalized Cut)



# Examples (texture classification with point-wise graph)



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## Graph-cuts

#### Bibliography

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# Full scene labeling (scene parsing)



Figure from Farabet et al , PAMI Tenenbaum and Barrow (1977)

- Segmentation in regions
- Building of the Region Adjacency Graph
- Labeling using a set of rules (expert system) :

1. on objects (size, color, texture,...)

on contextual relationships between objects (above, inside, near ...)

Generalization with fuzzy attributed graphs

# Markovian labeling (random graphs)

$$E(l) = \sum_{i} \Phi(d_i, l_i) + \beta \sum_{ij} \Psi(l_i, l_j)$$

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- Low-level applications:
  - pixel graphs
  - segmentation, classification, restoration
- High-level applications:
  - graph of super-pixels (SLIC, watershed, ...)
  - graph of primitives (edges, key-points, lines,...)
- CRF (Conditional Random Field) / MRF (Markov Random Field):
  - MRF:  $\Psi$  does not depend on *d* ("pure" prior)
  - CRF:  $\Psi$  depends on *d* (usually based on image gradient values)
  - $\Rightarrow$  pattern recognition, full scene labeling

# Example on a 3D RAG (T. Gud)

nuclei segmentation





3D anisotropic gradient



3D morphological closing



3D watershee



**3D** over-segmentation



result of graph labeling

#### Markovian relaxation



#### Example on a line graph



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# Example on a region adjacency graph



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### MRF and graph-cut optimization

Binary labeling (Greig et al. 89) :

$$\mathcal{E}(l) = \sum_i \Phi(d_i|l_i) + \sum_{(i,j)} eta(l_i - l_j)^2$$

efficient way of finding the global minimum by min-cut search

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#### MRF and graph-cut optimization



#### FIGURE 2.5

(a) The graph for binary MRF minimization. The edges in the cut are depicted as thick arrows. Each node other than  $v_s$  and  $v_t$  corresponds to a site. If a cut (\$, T) places a node in \$, the corresponding site is labeled 0; if it is in T, the site is labeled 1. The 0's and 1's at the bottom indicate the label each site is assigned. Here, the sites are arranged in 1D; but according to the neighborhood structure this can be any dimension as shown in (b) and (c).

(figure from "Image processing and analysis with graphs", Lezoray - Grady)

# MRF/CRF and graph-cut optimization Multi-level labeling (Boykov, Veksler) :

 $\Rightarrow$  generalization of the binary labeling Definition of two space moves (to go back to the binary labeling)

- $\alpha$ -expansion : source S and sink P correspond to label  $\alpha$  and the current label  $\overline{\alpha}$  ( $\Psi$  should be a metric)
- $\alpha \beta$  swap: source S for  $\alpha$  and sink P for  $\beta$  ( $\Psi$  should be a semi-metric)

Optimization by iterative mincut search:

- graph: nodes for super-pixels
- weights: depending on the current labeling
- good trade off time / efficiency compared to simulated annealing or ICM

But for multi-labeling no garantee on optimality of the solution  $(a,b) \in \mathbb{R}^{n}$ 

Interactive segmentation: "hard" constraints

Principle Background and object manually defined

 $\Rightarrow$  finding of a binary labeling minimizing an energy including "hard" constraints

Method Mincut search and edges with high weights (should not be cut)

#### Advantages

- easy introduction of "hard" constraints
- the manually defined areas permit to do a fast learning
- iterative algorithm

### Illustrations



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#### Interactive methods with mincut

#### Grab-cut

- take into account color
- two labels (background and object but with a Gaussian Mixture Model)
- CRF (conditional random field): regularization term weighted by the image gradient

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 iterative semi-supervised learning of the GMM parameters (after manual initialization and after each cut)

### Illustrations -GrabCut-



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# Deep learning and graph labeling for full scene labeling



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### Deep learning and graph labeling for full scene labeling



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#### Pattern recognition

- Object: defined by a set of primitives (nodes of the graph)
- Binary relationship of compatibility between nodes (edges of the graph)
- Clique: sub-set of primitives all compatible between each other = possible object configuration

recognition by maximal clique detection

#### Search of maximal cliques :

- NP-hard problem
- Building of a decision tree: a node of the tree = 1 clique of the graph
- pruning of the tree to suppress already found cliques
- ► Theorem: let S be a node of the search tree T, and let x be the first unexplored child of S to be explored. If all the sub-trees of S ∪ {x} have been generated, only the sons S not adjacent to x have to be explored.

### Example:maximal clique search



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# Example: buiding reconstruction by the maximal clique search (IGN) $% \left( IGN\right) =0$



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#### Graph matching

#### Correspondance problem:

- ► [Graph(s) of the model (atlas, map, model of object)
- Graph built from the data \_ bruite, --

Graph matching:

$$G = (X, E, \mu, \nu) \quad \rightarrow ? \quad G' = (X', E', \mu', \nu')$$

Graph isomorphism: bijective function  $f : X \to X'$ 

Too strict  $\Rightarrow$  isomorphisms of sub-graphs

## Sub-graph isomorphisms

There exists a sub-graph S' of G' such that f is an isomorphism from G to S'



There exists a sub-graph S of G and a sub-graph S' of G' such that f is an isomorphism from S to S'.

#### Graph isomorphisms: searching the maximal clique

search of the maximal clique of the association graph

principle: building of the association graph

maximal clique: sub-graph isomorphism



#### Graph isomorphisms: searching the maximal clique

search of the maximal clique of the association graph

- principle: building of the association graph
- maximal clique: sub-graph isomorphism



Sub-graph isomorphism: Ullman algorithm

- Principle : extension of the association set (v<sub>i</sub>, w<sub>x<sub>i</sub></sub>) until the G graph has been fully explored. In case of failure, go back in the association graph ("backtrack").
  Acceleration: "forward checking" before adding an association.
- Algorithm:
  - matrix of node associations
  - matrix of future possible associations for a given set of associations matrice
  - list of updated associations by "Backtrack" et "ForwardChecking"
- Complexity : worst case O(m<sup>n</sup>n<sup>2</sup>) (n ordre de X, m de X', n < m)</p>

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#### Error tolerant graph-matching

- Real world: noisy graphs, incomplete graphs, distorsions
- Distance between graphs (editing, cost function,...)
- Sub-graph isomorphism with error tolerance: search of the sub-graph G' with the minimum distance to G

- Optimal algorithms: A\*
- Approximate matching: genetic algorithms, simulated annealing, neural networks, probablistic relaxation,...
  - iterative minimistion of an objective function
  - better adapted for big graphs
  - problem of convergence and local minima

## Decomposition in common sub-graphs



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### Example

3D reconstruction by graph matching between a graph (data) and a library of model graphs (IGN)





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### Example - building reconstruction



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#### Example - building reconstruction Model graph and data graph matching



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### Matching with geometric transformation

- Graph = representation of the spatial information
- Matching = computation of the geometric transformation
  - polynomial deformation
  - elastic transformation (morphing)
- Matching approaches :
  - translation: maximum of correlation
  - Hough transform (in the parameter space)
  - RANSAC method: select randomly a set of matching points, compute the transformation, compute the score (depends on the number of matched pairs for the transformation)
  - AC-RANSAC: RANSAC + a contrario framework reducing the number of parameters (NFA to be set)

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## Example - MAC-RANSAC (PhD Julien Rabin)





(a) Paire d'images analysée.



(b) Reconnaissance de chacun des objets superposés.

## Example - MAC-RANSAC (PhD Julien Rabin)





(a) Paire d'images utilisée





#### Inexact matching

Optimization of a cost function

Dissimilarity cost beween nodes

$$c_N(a_D, a_M) = \sum \alpha_i d(a_i^N(a_D), a_i^N(a_M)) \quad \sum \alpha_i = 1$$

Dissimilarity cost between edges

$$C_{E}((a_{D}^{1}, a_{D}^{2}), (a_{M}^{1}, a_{M}^{2})) = \sum \beta_{j} d(a_{j}^{A}(a_{D}^{1}, a_{D}^{2}), a_{j}^{A}(a_{M}^{1}, a_{M}^{2})) \sum \beta_{j} = \beta_{j} d(a_{j}^{A}(a_{D}^{1}, a_{D}^{2}), a_{j}^{A}(a_{M}^{1}, a_{M}^{2}))$$

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Matching cost function h :

$$f(h) = \frac{\alpha}{|N_D|} \sum_{a_D \in N_D} c_N(a_D, h(a_D)) + \frac{1 - \alpha}{|E_D|} \sum_{\substack{(a_D^1, a_D^2) \in E_D}} c_E((a_D^1, a_D^2), (h(a_D^1), h(a_D^2)))$$

Optimization methods:

Tree search

- Expectation Maximization
- Genetic algorithms

## Example: brain structures (A. Perchant)



## Example : face structures (R. Cesar et al.)

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## Spectral method for graph matching (1)

Optimization of a cost function

- weighted adjacency matrix M
- nodes = potential assignments a = (i, i') (can be selected by descriptor matching)
- edges = M(a, b) agreement between the pairwise matchings a and b (geometric constraints)
- correspondance problem = finding a cluster C of assignments maximizing the inter-cluster score  $S = \sum_{a,b\in C} M(a,b)$  with additional constraints
- cluster C = vector x (with x(a) = 1 if  $a \in C$  and 0 else)

$$S = \sum_{a,b\in C} M(a,b) = x^T M x$$

$$x^* = argmax(x^T M x)$$

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+ constraints (one to one mapping)

Spectral method for graph matching (2)

#### Search of the optimal cluster

- number of assigments
- inter-connection between the assignments
- weights of the assignment

Spectral method: relaxation of the constraints on x

$$x^* = principal eigenvector(x^T M x)$$

+ introduction of the one-to-one correspondance constraints (iterative selection of  $a^* = argmax_{a \in L}(x^*(a))$ and suppression in  $x^*$  of the incompatible assignments)

## Example: point matching (Leordeanu, Hebert)



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 $\begin{aligned} d_{ab} &= \frac{d_{ij}+q}{d_{i'j'}+q} \\ \alpha_{ab} &= angle \text{ between the matchings} \\ \text{(with centring and normalization)} \\ M(a,b) &= (1-\gamma)c_{\alpha} + \gamma c_{d} \end{aligned}$ 

## Example:feature matching (Leordeanu, Hebert)



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## Example: factorized graph matching (Zhou, de la Torre)



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#### Spatial reasoning in images



(a) Example image.



(b) Labeled image: The blue regions represent the sea, the red and orange represent ships or boats and the yellow regions represent the docks.





(c) Concept hierarchy  $T_C$  in the context of harbors.

(d) Conceptual graph representing the spatial organization of some elements of Figure 5.8(b).

# Spatial reasoning in images





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