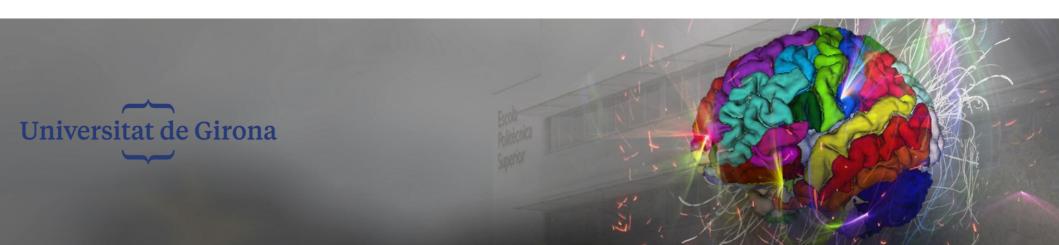


Computer Aided Diagnosis (CAD)

Deformable Models Segmentation





Part 1: Template matching & deformable template matching

Part 2: Active shape & Active appearance models

Part 3: Active contours & level sets





 Pattern matching is the act of checking a perceived sequence of tokens (or pixels, in computer vision) for the presence of the constituents of some pattern.

Pattern

Image

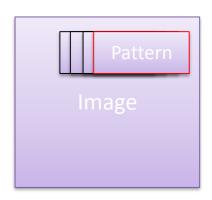
 In contrast to pattern recognition, the match usually has to be exact.

Pattern



 Template matching is a technique in digital image processing for finding small parts of an image which match a template image.

Pattern





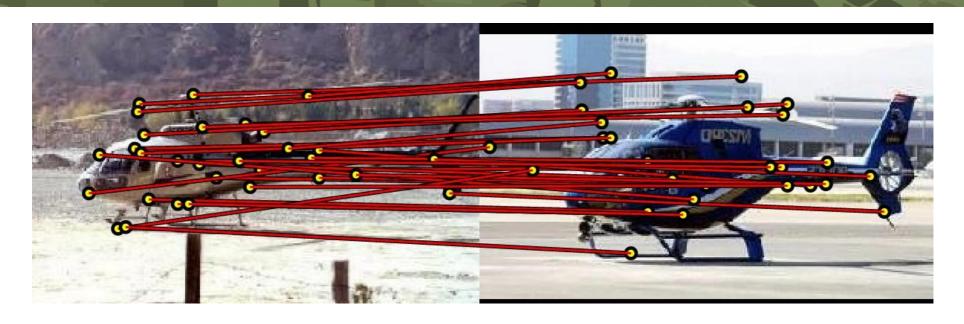


- Feature-based approach
 - If the template image has strong features
 - The approach may prove further useful if the match in the search image might be transformed in some fashion

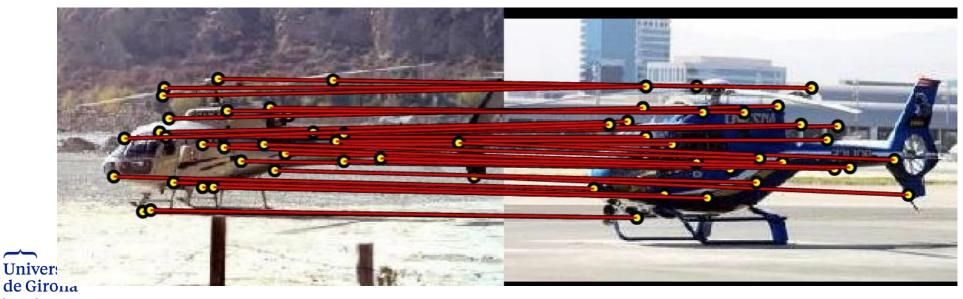








The deformable matching problem transforms to a correspondence problem

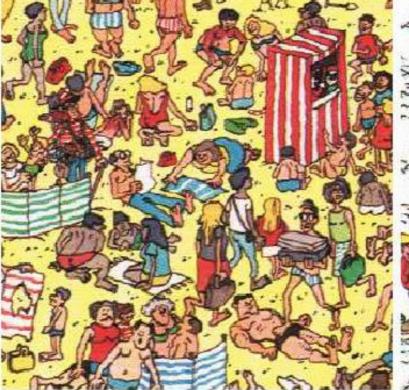




Template-based approach



Template







If you find it, matching!



If you find it, matching!



• To do: implement a simple algorithm for Wally detection in the image.





- To do: implement a simple algorithm for Wally detection in the image.
- Sliding window (borders?)
- Similarity definition:
 - Difference between images
 - Correlation
 - Entropy
 - Mutual information

... what problems do you see?

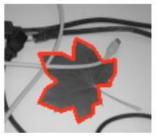




- Template matching from a template: which are the problems?
 - 2D rotation => add rotation as a new parameter
 - Scale => add scale as a new parameter
 - 3D rotation / perspective / orientation => we need a new view of the object or a 3D representation
 - Deformations => we need to know the ways that the object can deform
 - Occlusions => use a different similarity measure (per parts)





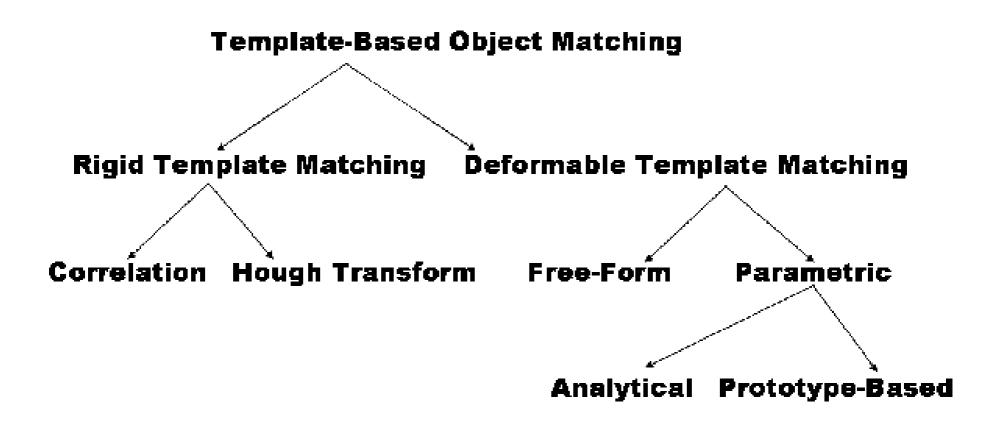








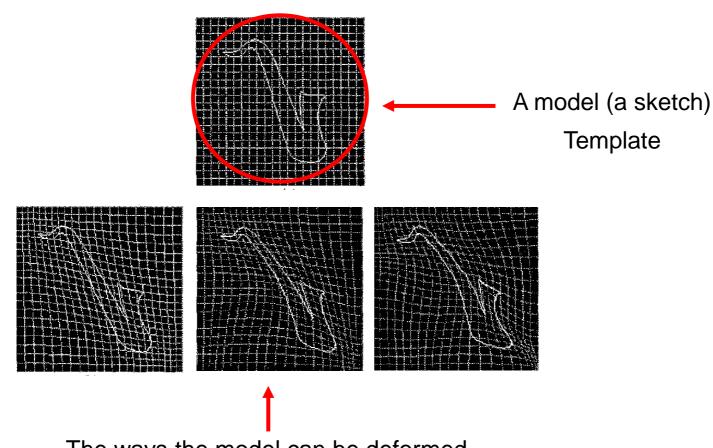








Prototype-based segmentation [Jain et al., 1996]

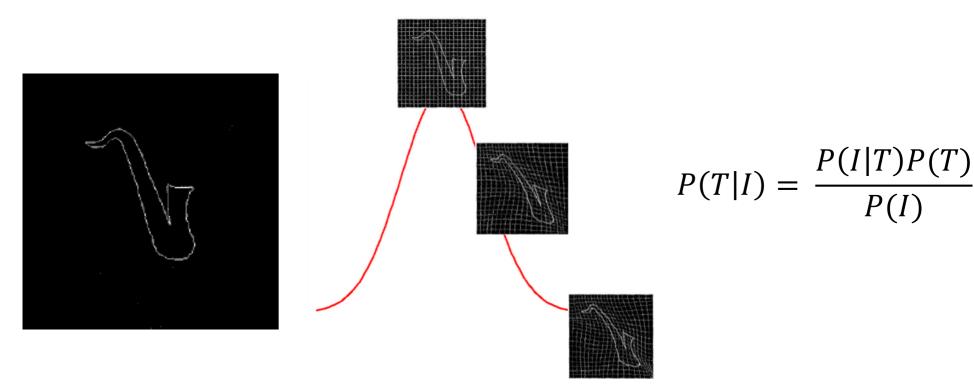




The ways the model can be deformed Deformations



 Uses prototype template, a set of deformation transformation, and a probabilistic model (Bayesian scheme) to find a match between the deformed template and objects in the image







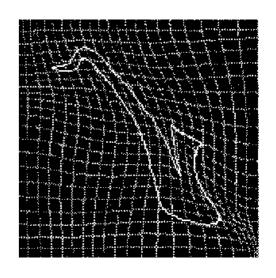
Main Idea: $(x, y) \rightarrow (x, y) + D_{\xi}(x, y)$

$$D_{\underline{\xi}}(x,y) = \sum_{m=1}^{M} \sum_{n=1}^{N} \frac{\xi_{mn}^{x} \cdot e_{mn}^{x} + \xi_{mn}^{y} \cdot e_{mn}^{y}}{\lambda_{mn}}$$

$$e_{mn}^{x}(x,y) = (2\sin(\pi nx)\cos(\pi my),0)$$

$$e_{mn}^{y}(x,y) = (0.2\cos(\pi mx)\sin(\pi ny))$$

$$\underline{\xi} = \{\xi_{mn}^x, \xi_{mn}^y, m, n\}$$

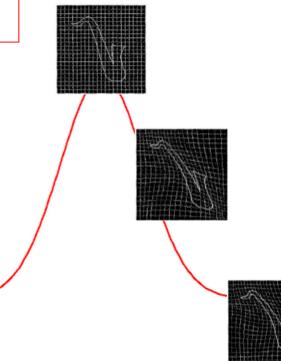






 Gaussian distribution to biased the possible deformed templates.

$$P_r\left(\underline{\xi}\right) = \frac{1}{(2\pi\sigma^2)^{MN}} e^{-\frac{1}{2\sigma^2}\sum_{m,n}(\xi_{mn}^x)^2 + \xi_{mn}^y}$$

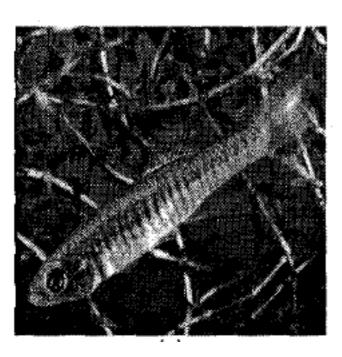






The likelihood has the image information:

$$P_r\left(I\middle|\mathbf{s},\theta,\underline{\xi},\underline{d}\right) = \alpha e^{-\frac{1}{n_t}\sum(1+\Phi(x,y)|\cos(\beta(x,y)))}$$
$$\Phi(x,y) = -\exp\left\{-\rho\left(\delta_x^2 + \delta_y^2\right)^{\frac{1}{2}}\right\},$$









- The Bayesian schema is employed to integrate the prior knowledge of the template and the observed object in the input image.
- We want to find the probability of observing the template (T) given the image (I):

$$P(T|I) = \frac{P(I|T)P(T)}{P(I)}$$

Maximising P(T|I) is equivalent to the minimise:

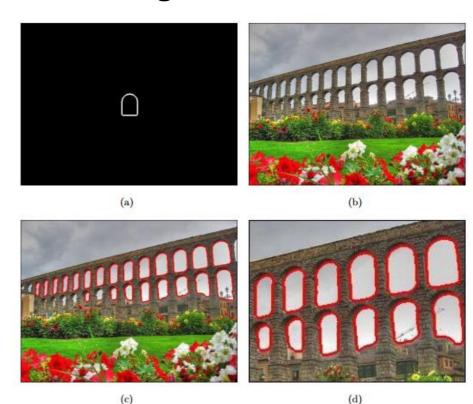
$$\mathcal{L}(\mathcal{T}_{s,\Theta,\underline{\xi},\underline{d}},Y) = \frac{M}{N} \sum_{i=1}^{N} (\pi x^{2})^{2}$$



$$\mathcal{E}\left(\mathcal{T}_{s,\Theta,\underline{\xi},\underline{d}},Y\right) + \gamma \sum_{m=1}^{M} \sum_{n=1}^{N} \left(\xi_{mn}^{x^{2}} + \xi_{mn}^{y^{2}}\right)$$



- Consist of a coarse-to-fine matching:
- First: roughly locate the global optima efficiently without regard to accuracy.
- The finer stage levels initialized using the good candidates screened from the previous stage.



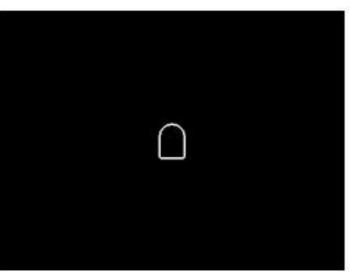




· Consist of a soarce to fine matching.

• Fir: re_§

• The scr

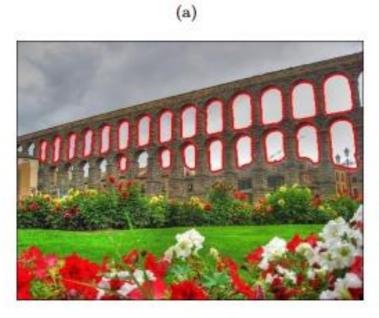




(b)

out

ates





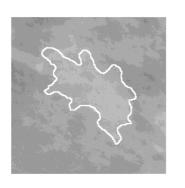


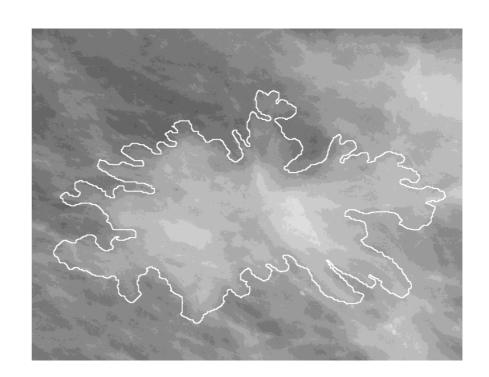
(d)



- How can we do this for masses?
 - There is a large range of mass shapes and sizes







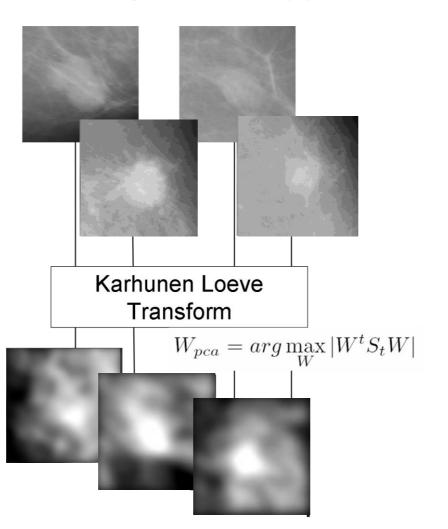


[Oliver PhD, 2007]

[Freixenet et al. 2008] Eigendetection of masses considering false positive reduction and breast density information. Medical Physics



We adapted the eigenfaces approach*



Initial mass database

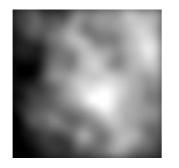
PCA transform

Eigenmasses





- Eigenfaces are calculated following these steps:
 - 1) Choose a (representative) training set
 - 2) Transform the images in a single column-vector
 - 3) Calculate the mean vector and subtract it to all vectors
 - 4) Calculate the covariance matrix
 - 5) Calculate its eigenvectors and eigenvalues
 - 6) Choose the principal components
 - 7) Optionally, re-arrange the principal vectors in images













Mathematically, the template is defined as:

$$\psi^{0}(x,y) = \frac{1}{N} \sum_{k=1}^{N} w_{k} W_{k}(x,y)$$

But we use only the contours of the eigenmasses

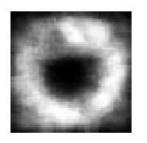
$$\nabla \psi^0(x,y) = \nabla \left\{ \frac{1}{N} \sum_{k=1}^N w_k W_k(x,y) \right\} =$$

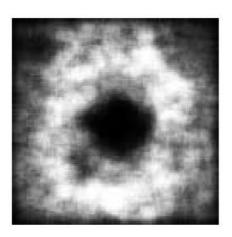
$$= \frac{1}{N} \sum_{k=1}^N w_k \nabla W_k(x,y)$$

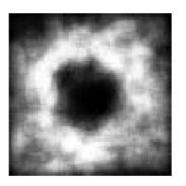


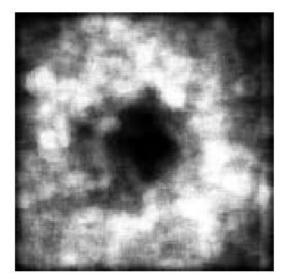


 We cluster the database according to a set of sizes. Thus, we have a model per size













The template is the model. But, what about the deformations?

$$\nabla \psi^d(x,y) = \kappa \sum_{k=1}^N \xi_k w_k \nabla W_k(x,y)$$

- We assume that masses are linear combinations of the eigenmasses. Hence, we introduced a vector of coefficients which modifies the template
- Once we have defined the template and its plausible deformations, we have to perform the matching step





To match the model into new mammograms we follow a probabilistic Bayesian strategy

$$posterior = \frac{prior*likelihood}{evidence}$$

The prior distribution is used to bias the global and local deformations

$$Pr(s, d, \xi) = K \exp\{-\frac{1}{2\sigma^2} \sum_{k=1}^{N} (\xi_k - 1)^2\}$$

 A deformed template with a geometric shape similar to the prototype template is favoured





 The likelihood is a measurement of the similarity between the deformed template and the object(s) present in the image

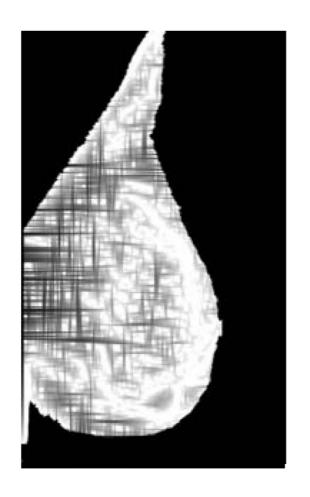
 As we have defined a contour-based template, we have to obtain a contour-enhanced mammogram representation

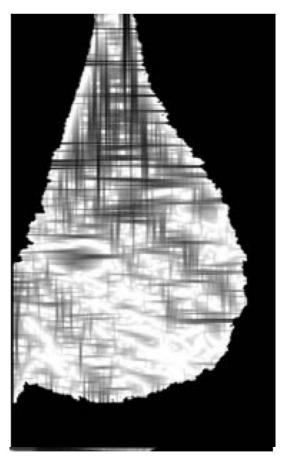
$$\phi_Y(x,y) = -\exp(-\rho\sqrt{\delta_x^2 + \delta_y^2})$$

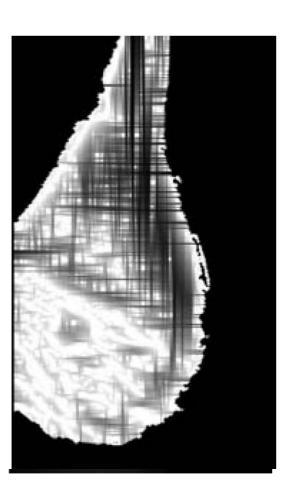




Potential images











Likelihood probability

$$Pr(Y|s,d,\xi) = \alpha \exp\{-\Upsilon(\psi^{s,\xi,d},Y)\}$$

Where

$$\Upsilon(\psi^{s,\xi,d}, Y) = \frac{1}{T} \sum_{x,y \in \psi^{s,\xi,d}} (1 + \phi_Y(x,y) | cos(\beta(x,y))|)$$

 This definition requires that the template boundary agrees with the image edges not only in position, but also in the tangent direction





The posterior probability is what we want

$$Pr(s, d, \xi|Y) = \frac{Pr(s, d, \xi) Pr(Y|s, d, \xi)}{Pr(Y)}$$

 Doing the operations we found that maximising the posterior was equivalent to minimise the following equation

$$\Lambda(\psi^{s,\xi,d}, Y) = \sum_{k=1}^{N} (\xi_k - 1)^2 + \Upsilon(\psi^{s,\xi,d}, Y)$$

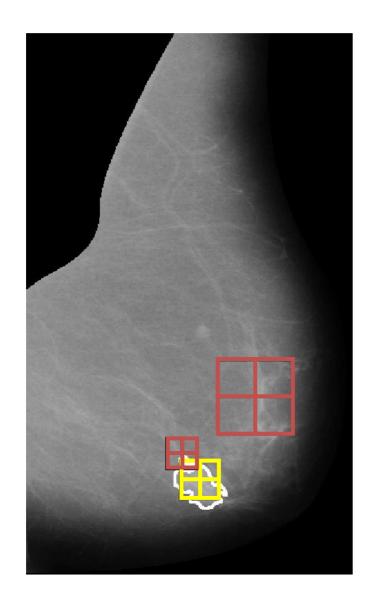
Related to the deviation of the templates from the prototype

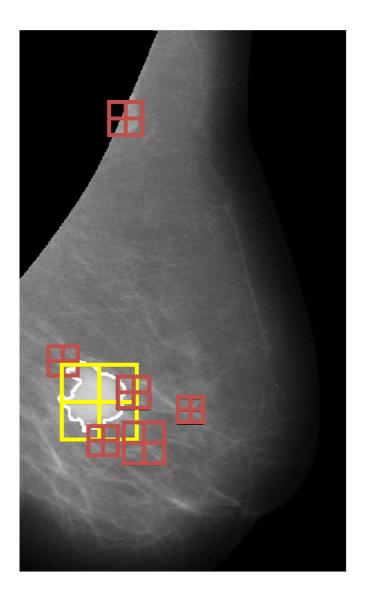
Related to the fitness of the deformed template to the boundaries





Examples of the algorithm (put a dense mammo):









Examples of the algorithm:





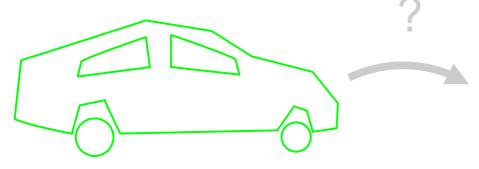




The modern approach is dividing the model by parts:

Image





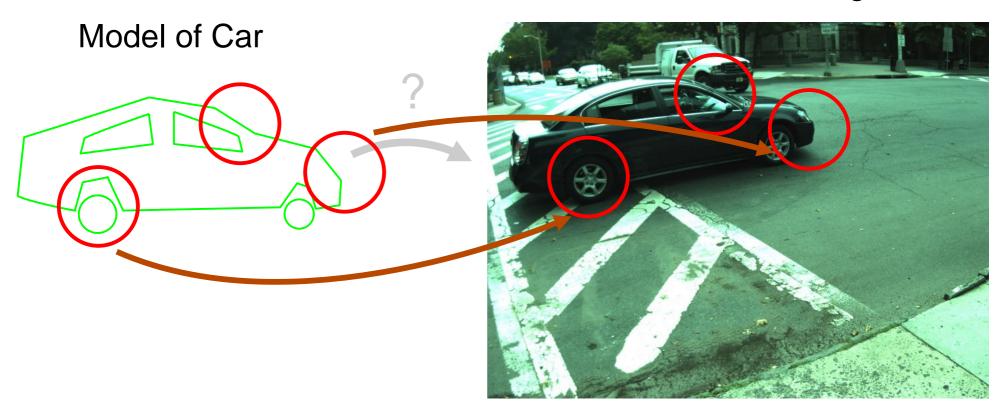






The modern approach is dividing the model by parts:

Image

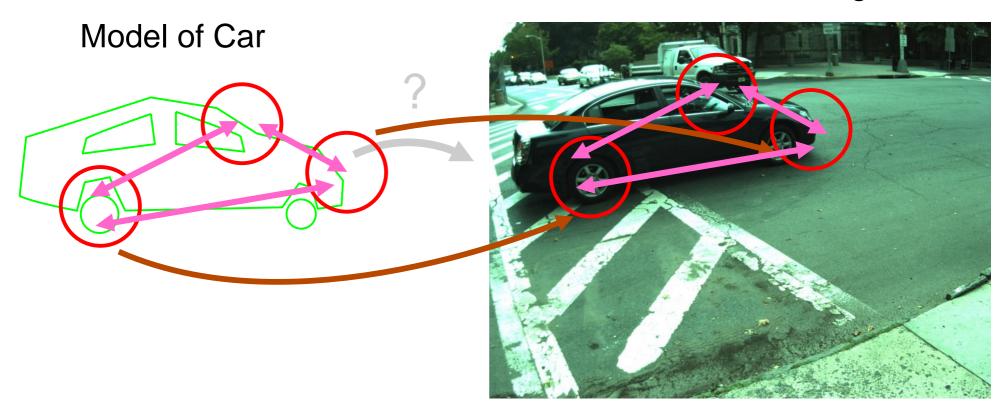






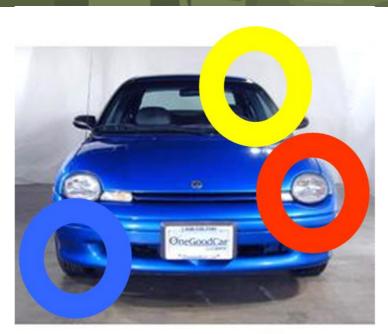
The modern approach is dividing the model by parts:

Image



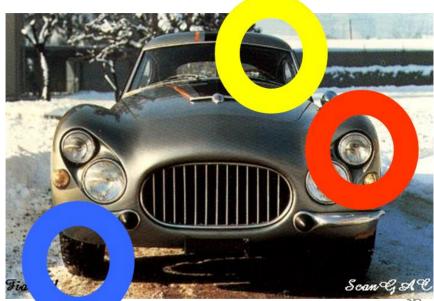
















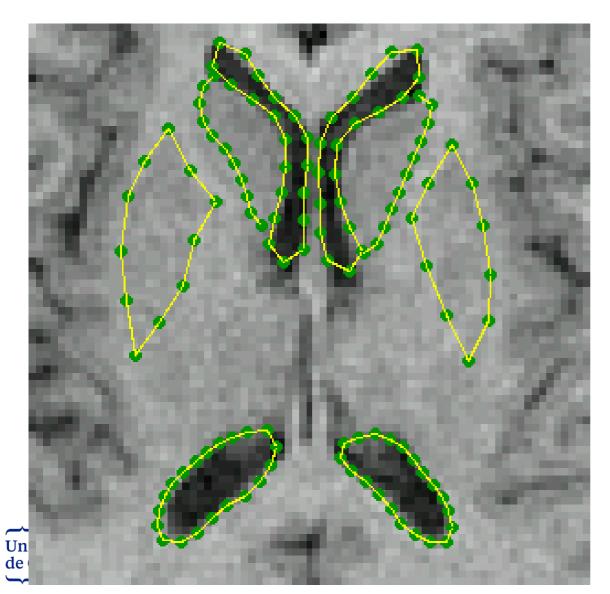
- A deformable template matching approach but using a different way for creating the model
- Developed by Tim Cootes, Active Shape Models-Their Training and Application, CVIU, 1995
- Posterior improvement: Active appearance models, PAMI,
 2001

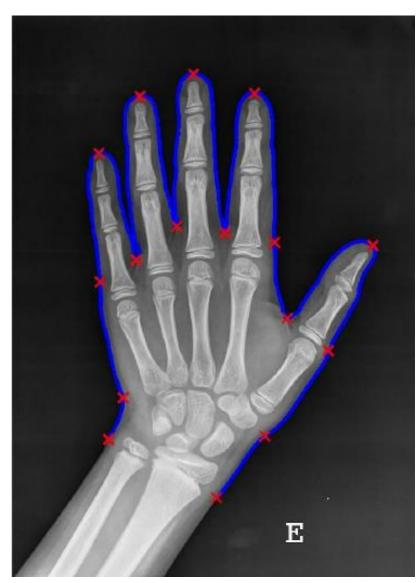
Widely used in medical imaging





Widely used in medical imaging





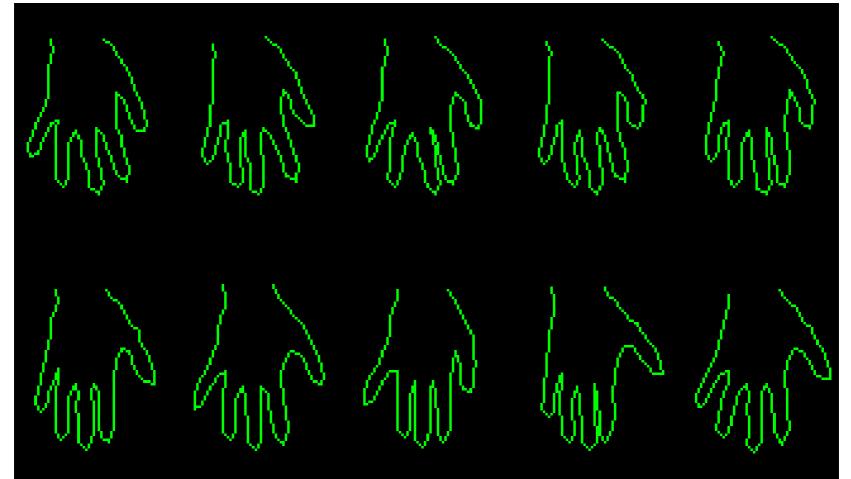


- Active Shape models are based in mainly two steps:
 - Build a statistical shape model (PDM: point distribution model)
 - Adjust it to the image
- The shape is represented by keypoints
- In all the examples the keypoints are placed in the same exact position





- Consider the outline of a hand, represented by 72 labelled points.
- Here are some examples from a training set:







- Mean Model alignment: Procrustes analysis
 - The Procrustes distance is a least-squares type shape metric that requires two aligned shapes with one-to-one point correspondence
 - The alignment part involves four steps:
 - Compute the centroid of each shape.
 - Re-scale each shape to have equal size.
 - Align w.r.t. position the two shapes at their centroids.
 - Align w.r.t. orientation by rotation.





- Statistical Models: initial image processing
 - Procrustes analysis for 2 vectors: find transformation which minimizes

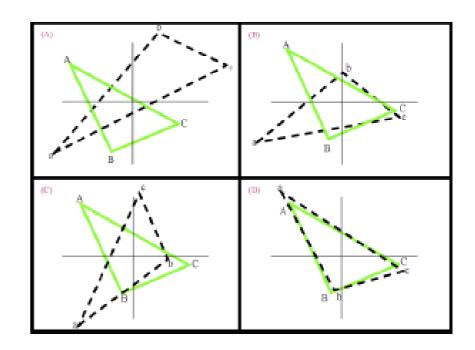
$$|x_1 - T(x_2)|^2$$

– for n vectors, minimize:

$$\sum |m - Ti(x_i)|^2$$

$$\mathbf{m} = \frac{1}{n} \sum T_i(\mathbf{x}_i) \quad |\mathbf{m}| = 1$$

- Resulting shapes have
 - Identical centre of gravity
 - Approximately the same scale and orientation





Each hand is represented by a 2n element vector

$$x = (x_1, ..., x_n, y_1, ..., y_n)$$

- We use Principal Component Analysis (PCA) to pick out the main axes of the cloud, and model only the first few, which account for the majority of the variation.
- The shape model is then

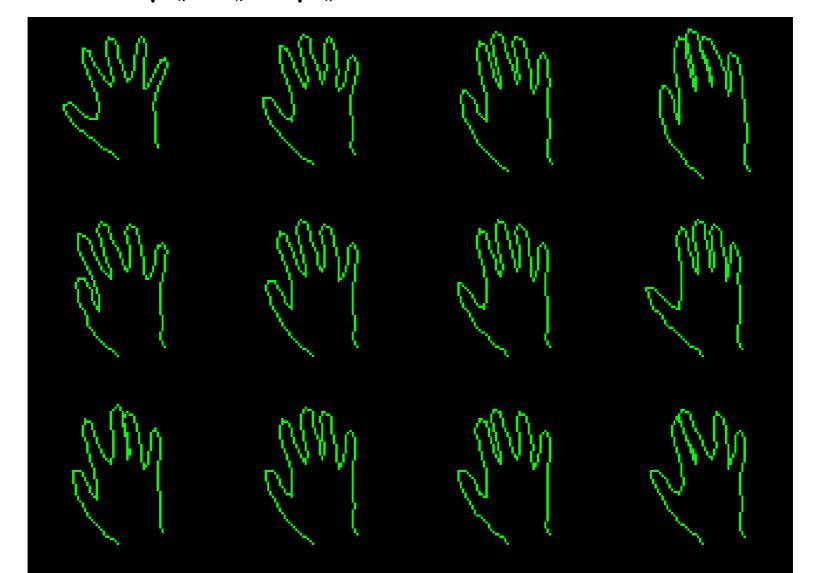
$$x = x_mean + Pb$$

- where
 - x_mean is the mean of the aligned training examples,
 - P is a 2n x t matrix whose columns are unit vectors along the principal axes of the cloud (the t eigenvectors), and
 - b is a t element vector of shape parameters.





• By varying the shape parameters (b) within limits learnt from the training set $(-3\sqrt{\lambda_k} \le b_k \le 3\sqrt{\lambda_k})$, we can generate new examples.





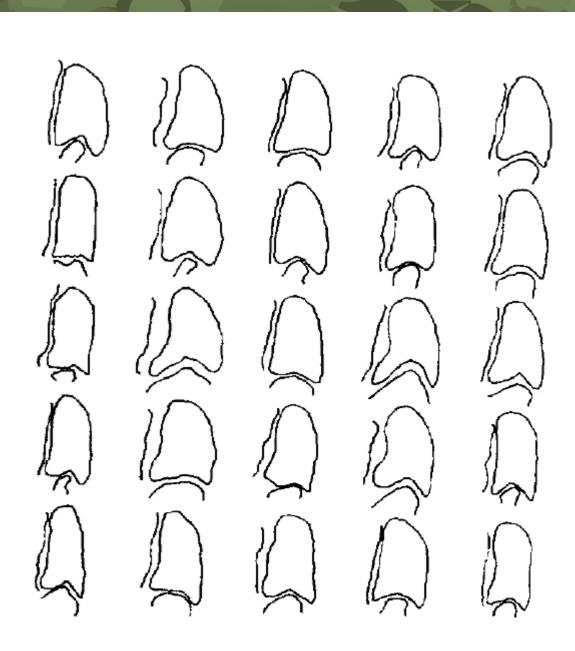


Heart Example

- 66 examples
- 96 points
 - Left ventricle
 - Right ventricle
 - Left atrium
- Traced by cardiologists

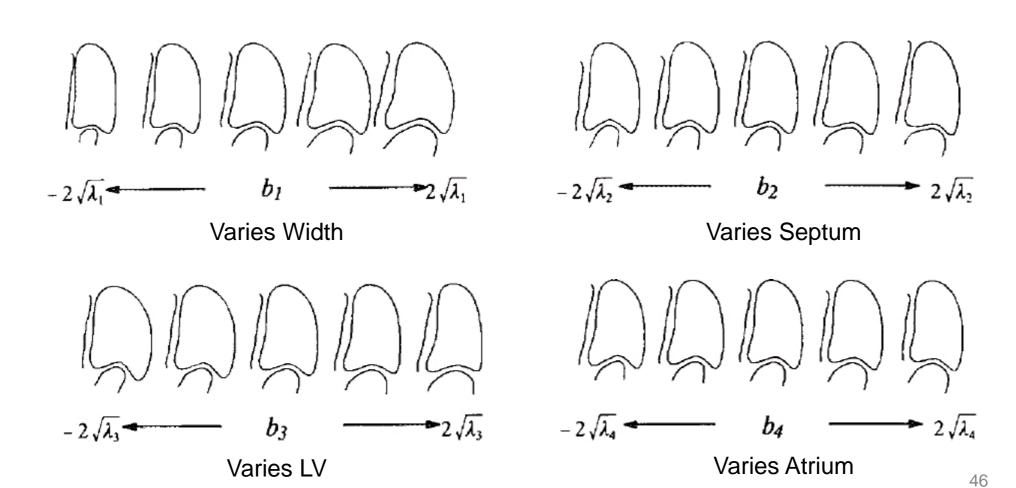
Eigenvalues of the Covariance Matrix Derived from a Set of Heart Ventricle Shapes

	Eigenvalue	$\frac{\lambda_i}{\lambda_T} \times 100\%$
	λ,	37%
	λ_2	17%
	λ_3	13%
	λ_4	7%
<u></u>	λ_5	6%
Universit		4%





Heart Example





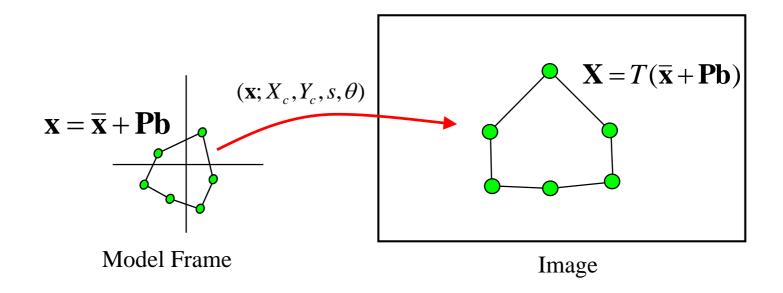


- How to adjust to a new image?
 - Put the mean model over the image
 - Look along normals through each model point to find the best local match for the model of the image appearance at that point (eg strongest nearby edge)
 - Update the pose and shape parameters to best fit the model instance to the found points
 - Repeat steps 2 & 3 until convergence
- The performance can be significantly improved using a multi-resolution implementation





- Active Shape Models (ASMs)
 - Must apply global transformation ${f T}$ to place in image.

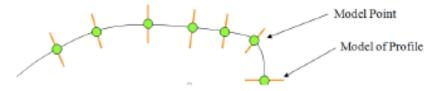


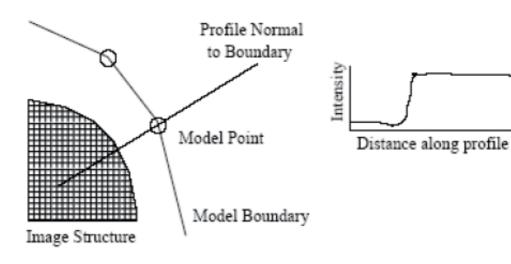
where (X_c, Y_c) is the position of the centre of the model in the image frame, rotated by θ and scaled by s.





- Active Shape Models (ASMs): how it works
 - Match shape model to new image
 - Require:
 - Statistical shape model
 - Model of image structure at each po

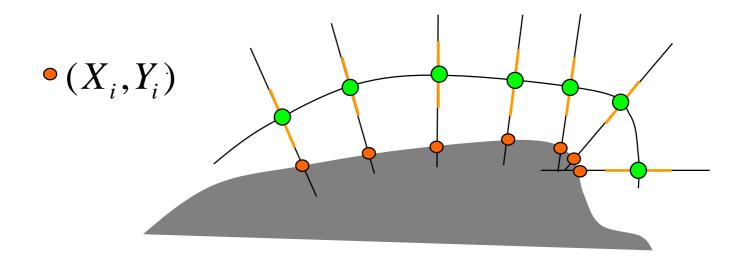








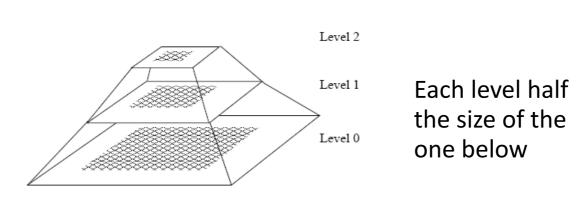
- Active Shape Models (ASMs): how it works
 - Algorithm
 - Examine a region of the image around each point (X_i, Y_i) to find the best nearby match.
 - Update parameters (X_t, Y_t, s, θ, b) to best fit the new found points.
 - Repeat until convergence.







- Active Shape Models (ASMs): how it works
 - Improved efficiency
 - Multi-resolution framework: gaussian image pyramid formed by repeated smoothing and sub-sampling.



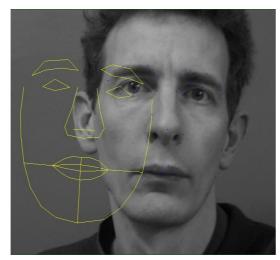


- Rapid location of the boundary of objects with similar shape
- Useful to classify objects based on shape or appearance
- Useful when approximate location of target object is known

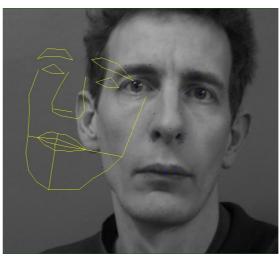




- Active Shape Models (ASMs): Limitations
 - Can fail if the initial guess is too far from the target.
 - Problems when position/size/orientation of targets is not known approximately.
 - Doesn't work with widely varying shapes.
 - The model can only deform in ways observed in the training set.
 If is not there, the model will not fit to it.







Search result



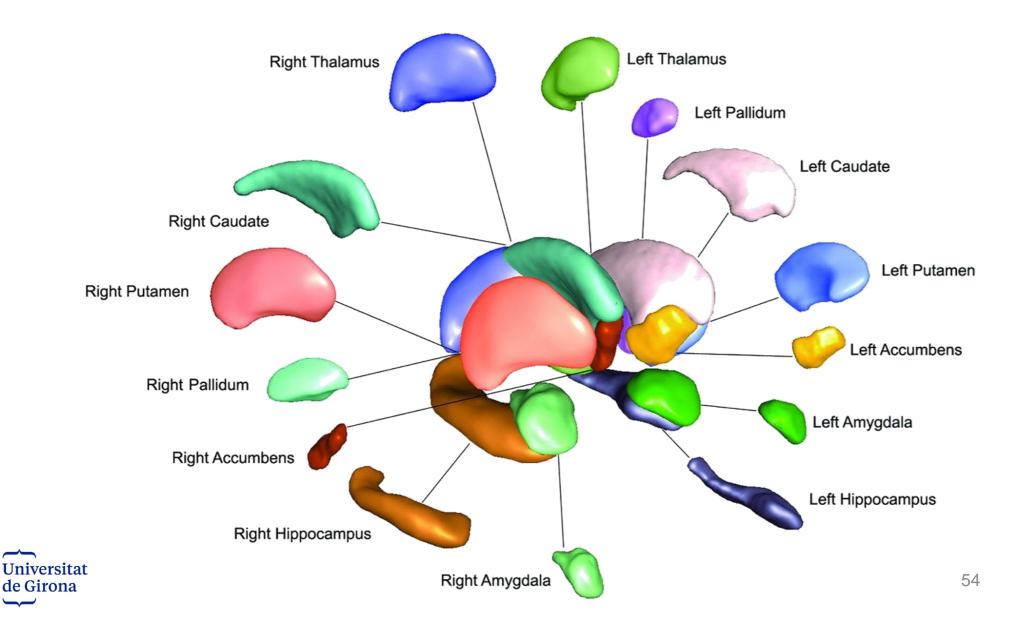


- The Active Appearance Model (AAM) is a generalization of the Active Shape Model approach.
- AAM uses all the information in the image region covered by the target object, rather than just near modelled edges.
- Widely used in medical imaging:
- Examples: brain structure segmentation, prostate segmentation, ...



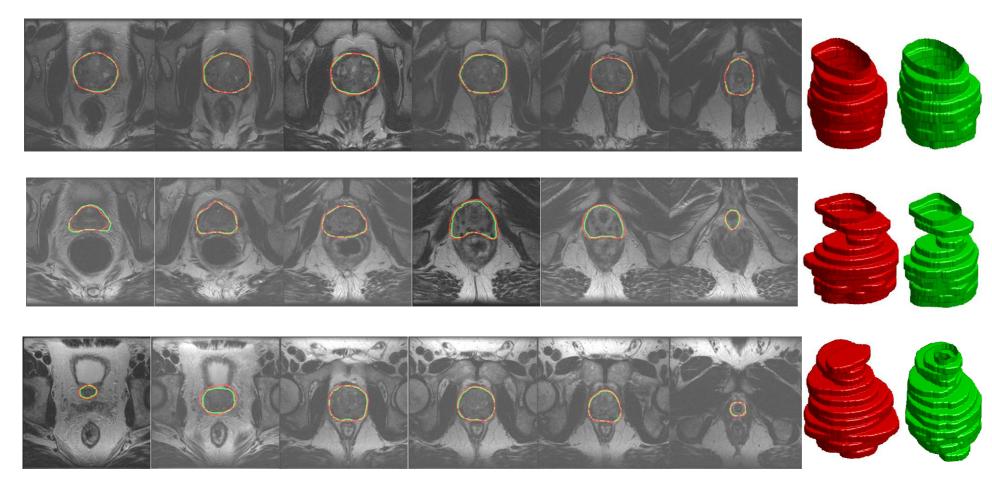


FIRST: brain structures segmentation





 Ghose PhD, 2013: active appearance model for prostate segmentation







 Method: given a set of training images, labelled with landmark points, we can use image warping to deform each image so that the object has the mean shape, then build a statistical model of the grey-levels across the object.

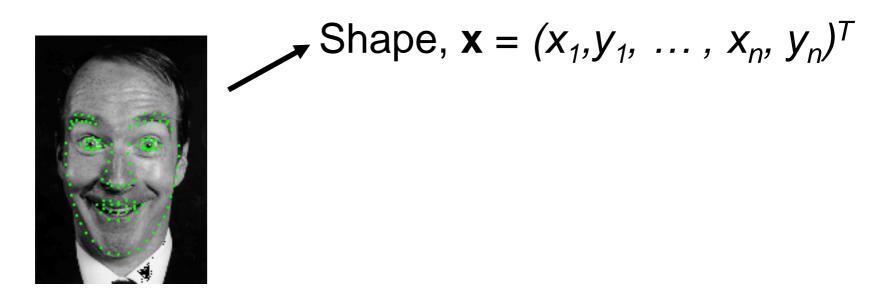
Mainly:

- Shape models represent shape variation
- Eigen-models can represent texture variation
- But... combined appearance models represent both!





For each example extract shape vector



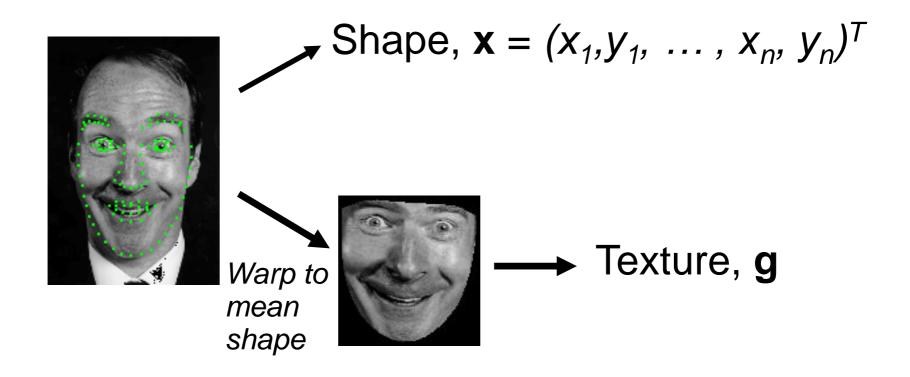
Build statistical shape model,

$$\mathbf{x} = \overline{\mathbf{x}} + \mathbf{P}_{s} \mathbf{b}_{s}$$





• For each example, extract texture vector

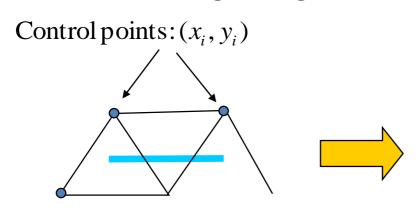


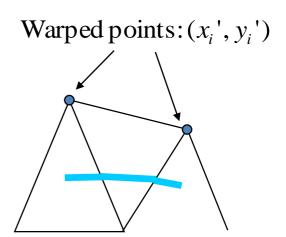
 Given corresponding points in two images, how do we warp one into the other?



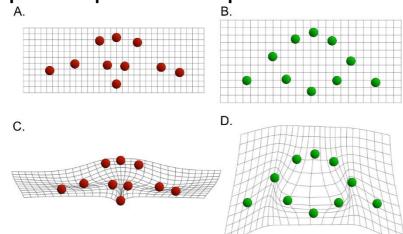


- Warping texture:
- Two common solutions
 - Piece-wise linear using triangle mesh





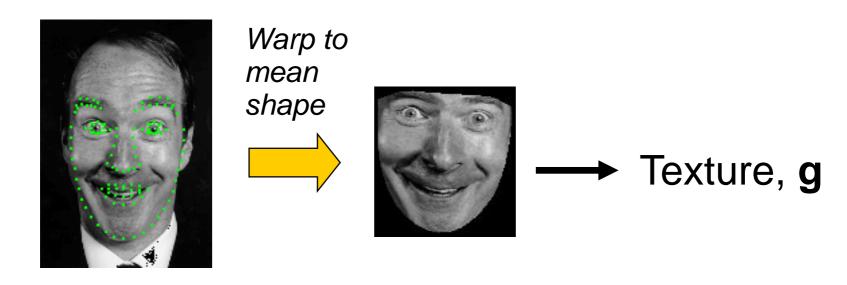
Thin-plate spline interpolation







For each example, extract texture vector



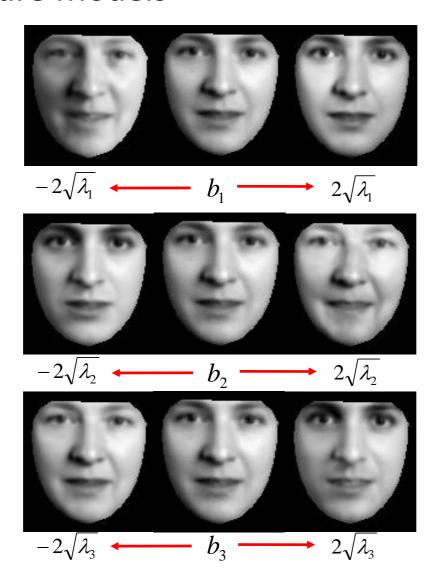
- Normalise vectors (as for eigenfaces)
- Build eigen-model

$$\mathbf{g} = \overline{\mathbf{g}} + \mathbf{P}_g \mathbf{b}_g$$





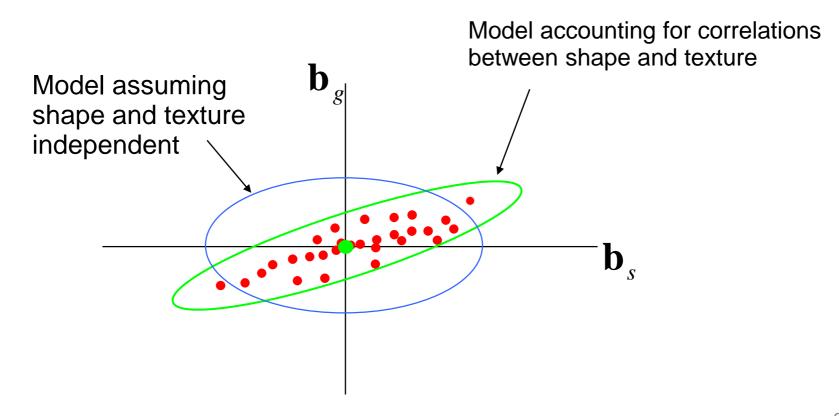
Face Texture Models







- Shape and texture often correlated:
 - When smile, shadows change (texture) and shape changes
- Learning this correlation leads to more compact (and specific) model







- For each image in training set we have best fitting shape and texture parameters \mathbf{b}_s , \mathbf{b}_g
- How to decorrelate the parameters? PCA again!
- Given the following vector:

$$\mathbf{b}_{c} = \begin{pmatrix} \mathbf{W}\mathbf{b}_{s} \\ \mathbf{b}_{g} \end{pmatrix} \qquad \mathbf{x} = \overline{\mathbf{x}} + \mathbf{P}_{s}\mathbf{b}_{s} \\ \mathbf{g} = \overline{\mathbf{g}} + \mathbf{P}_{g}\mathbf{b}_{g}$$

apply PCA:

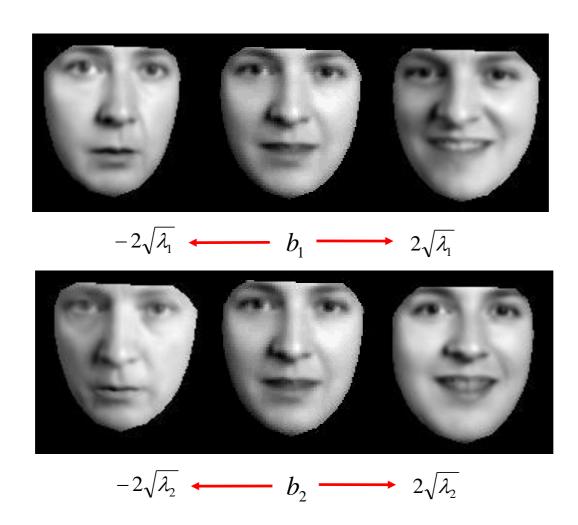
$$\mathbf{b}_{c} = \mathbf{Q}\mathbf{c} = \begin{pmatrix} \mathbf{Q}_{s} \\ \mathbf{Q}_{g} \end{pmatrix} \mathbf{c} \qquad \mathbf{x} = \overline{\mathbf{x}} + \mathbf{Q}_{s}\mathbf{c}$$
$$\mathbf{g} = \overline{\mathbf{g}} + \mathbf{Q}_{g}\mathbf{c}$$

W is a diagonal matrix to compensate the difference in units between shape and grey-level parameters

Hence, just varying c changes both shape and texture



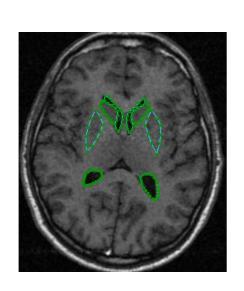


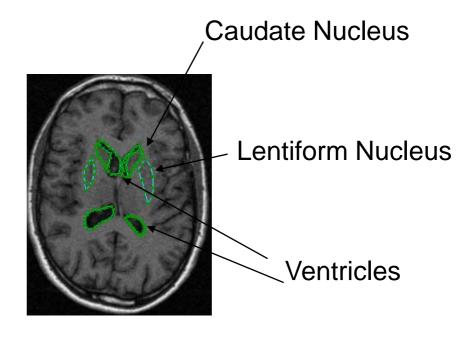






- Sub-cortical structures:
 - 72 examples
 - 123 points
 - 5000 pixel model



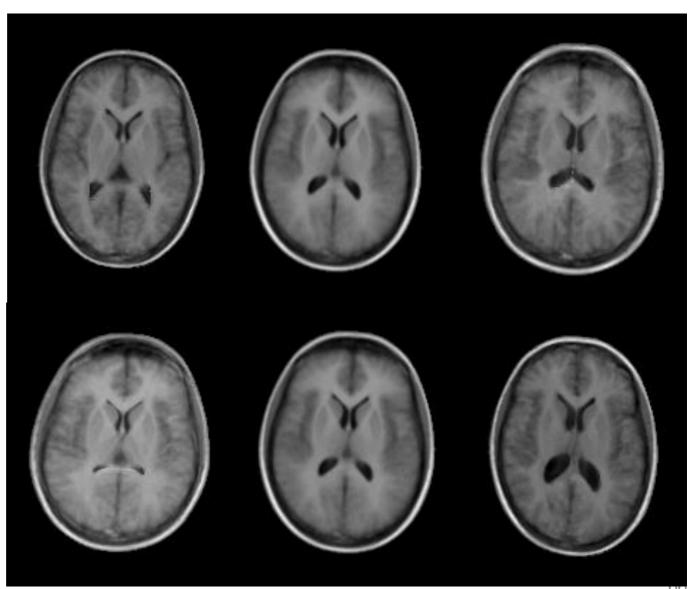






Combined Mode 1

Combined Mode 2







To know more...

- "Deformable template models: A review", A.K. Jain et al. Signal Processing, Vol 71, 109-129, 1996
- "Object Matching Using Deformable Templates", A.K. Jain et al. IEEE Pattern Analysis and Applications, Vol 18(2), 267-278, 1996
- "Active Shape Models Their Training and Application". T.F. Cootes et al. Computer Vision and Image Understanding Vol. 61(1), 38-59, 1995.
- "Active Appearance Models". T.F. Cootes et al. Proc. European Conference on Computer Vision 1998(2), pp. 484-498, 1998.
- "3-D Active Appearance Models: Segmentation of Cardiac MR and Ultrasound Images". S.C. Mitchell et al. IEEE Transactions on Medical Imaging, vol. 21(9), 1167-1178, 2002.

