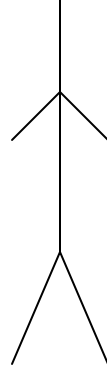
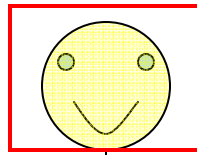
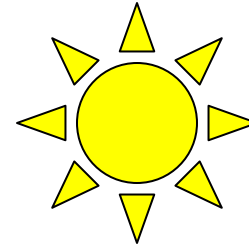


Geometric Pattern Matching

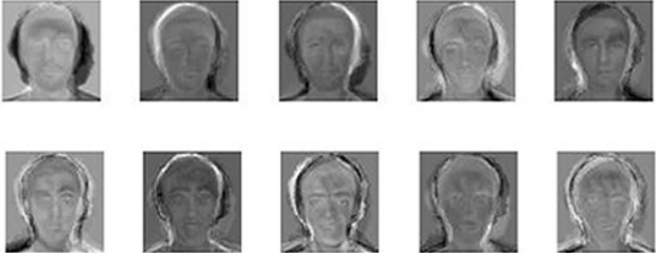
Piotr Indyk

Matching in a Scene



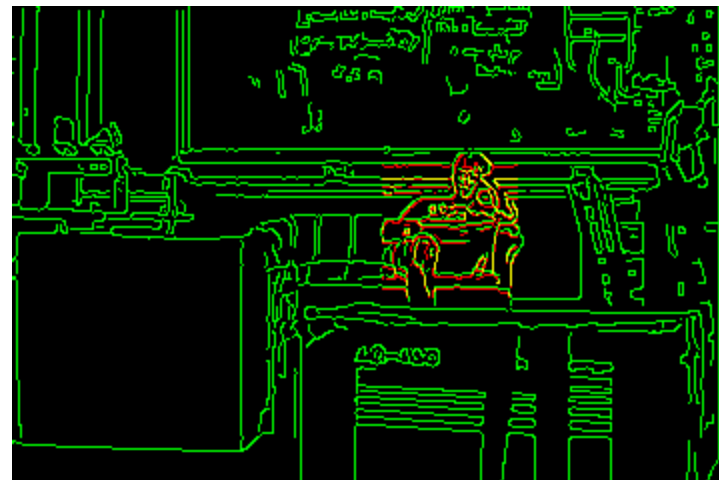
Eigenfaces



$$= 0.02 + 0.03 + 0.01 + \dots$$


Formalization: Shapes

- Today:
 - A shape is a set A of points in \mathbb{R}^2
 - $|A|=n$
- In general, A could consist of segments etc.



Formalization: (Dis)similarity

- Hausdorff distance
 - $DH(A,B) = \max_{a \in A} \min_{b \in B} \|a-b\|$
 - $H(A,B) = \max[DH(A,B), DH(B,A)]$
- Earth-Mover Distance
 - Minimum cost of a one-to-one matching between A and B
 - $EMD(A,B) = \min_{f:A \rightarrow B} \sum_{a \in A} \|a-f(a)\|$, f is 1:1
- Bottleneck matching
 - $BM(A,B) = \min_{f:A \rightarrow B} \max_{a \in A} \|a-f(a)\|$, f is 1:1

Alignment

- In general, **A** and **B** are not **aligned**
- So, in general, we want
 - $DH_T(A,B) = \min_{t \in T} DH(t(A), B)$, where
 - T=translations
 - T=translations and rotations
 - ...
 - Same for **H**

Algorithms

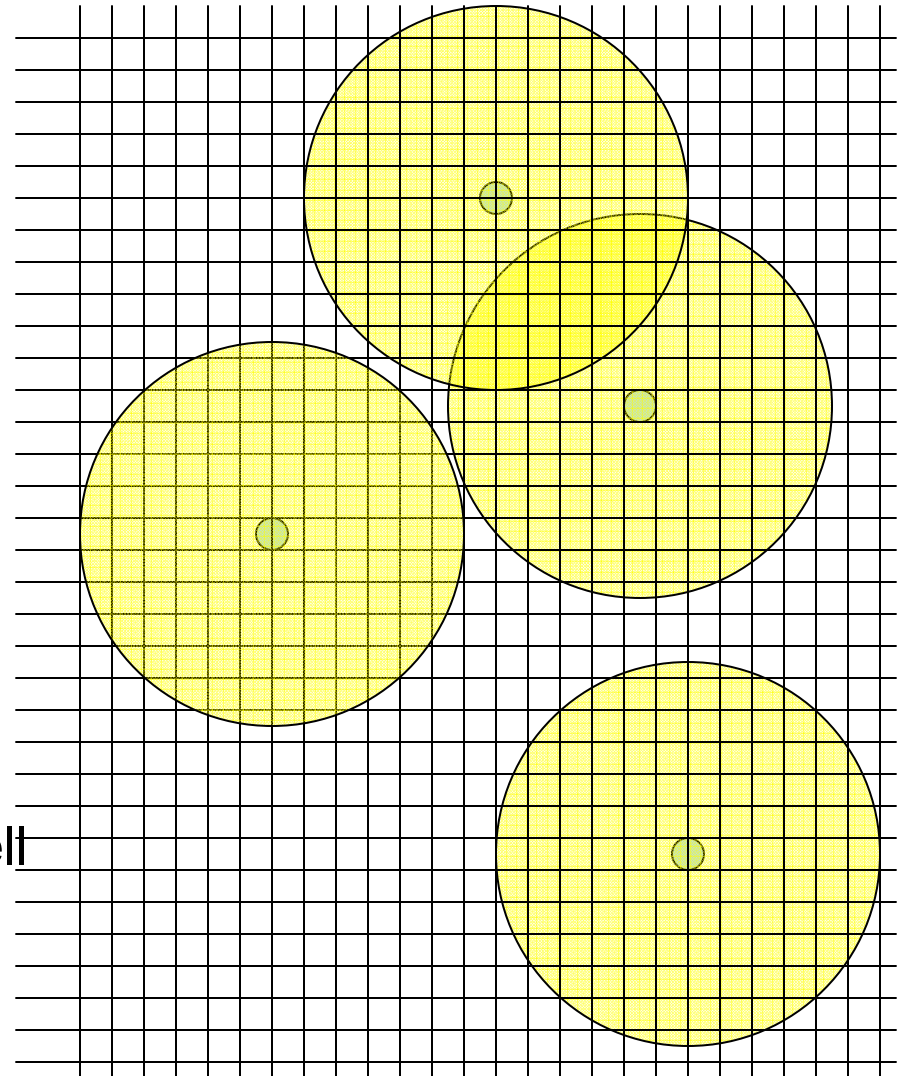
- Exact DH $O(n \log n)$
- Approx. DH $O(n)$
- Exact DH_T $O(n^4)$
- Approx. DH_T $O(n^2)$
- Approx. H_T $O(n \log n)$
- $DH_{T+R}=0$ $O()$

Computing $H(A,B)$

- Given A, B , how fast can we compute $H(A,B)$?
 - We will compute $DH(A,B)$
 - Construct a Voronoi diagram V for B
 - Construct a point location structure for V
 - For each a in A , find its NN in B
- Total time: $O(n \log n)$

Approximate Hausdorff

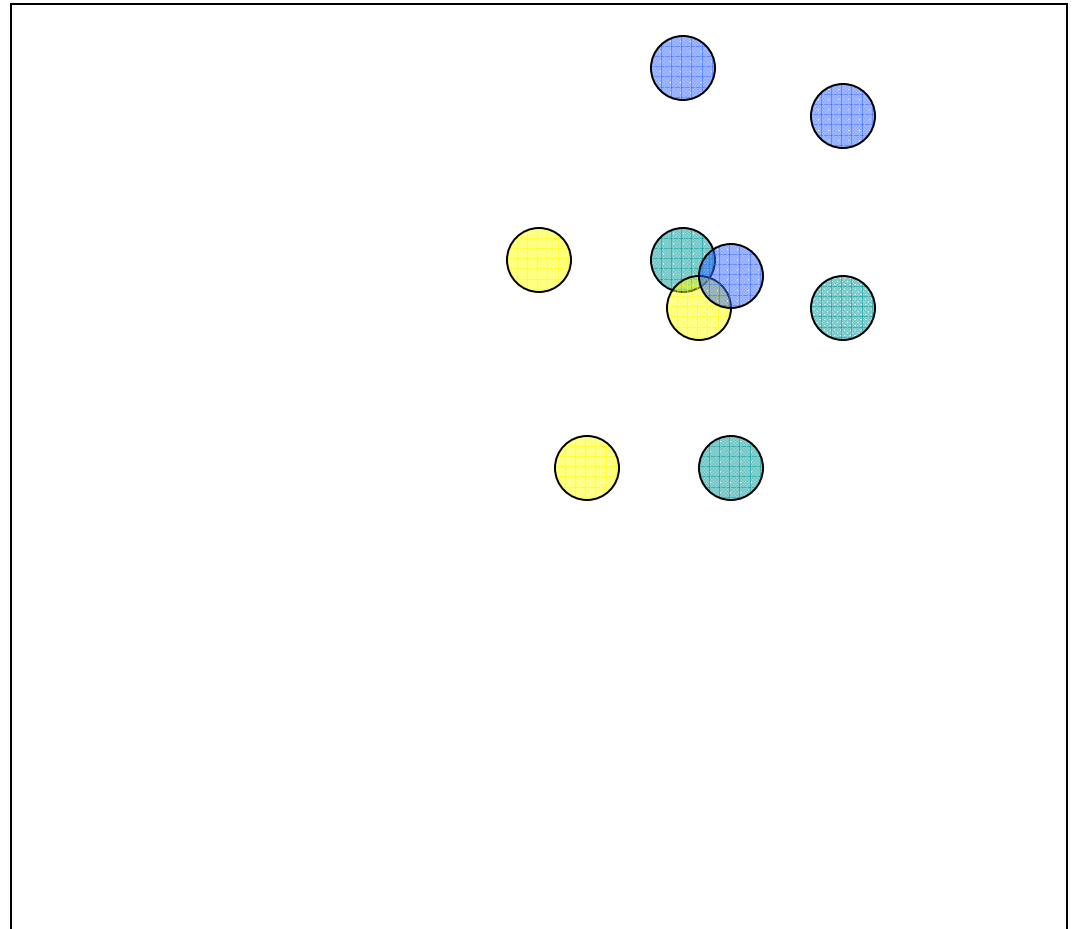
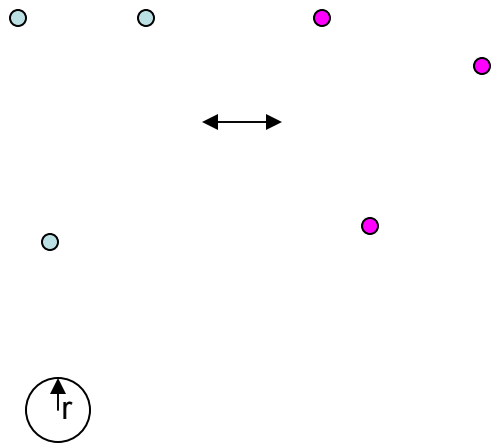
- Assume we just want an algorithm that:
 - If $DH(A,B) \leq r$, answers YES
 - If $DH(A,B) \geq (1 + \epsilon)r$, answers NO
- Algorithm:
 - Impose a grid with cell diameter ϵr
 - For each $b \in B$, mark all cells within distance r from b
 - For each $a \in A$, check if a 's cell is marked
- Time: $O(n/\epsilon^2)$



Decision Problem

- Again, focus on if $DH_T(A,B) \leq r$
- For $a \in A$, define
$$T(a) = \{ t : \exists b \in B \ ||t(a)-b|| \leq r \}$$
- $DH_T(A,B) \leq r$ iff $\bigcap_{a \in A} T(a)$ is non-empty

Example



Algorithm

- Compute the arrangement of all disks
- Check for non-emptiness
- Analysis:
 - Number of disks: n^2
 - Size of the arrangement: $O(n^4)$
 - Can compute the arrangement in time $O(n^4)$
 - Total time: $O(n^4)$
- Can be improved to $O(n^3 \log n)$
[Chew, Goodrich, Huttenlocher, Kedem, Kleinberg, Kravets'93]

Running time

- Running time pretty high
- Can we speed it up?
- A $(1+\varepsilon)$ -approximate algorithm for the decision version of DH_T :
 - Impose a grid with pixel diameter εr
 - Approximate each disk by $O(1/\varepsilon^2)$ cells
 - Each $T(a)$ is approximated by $O(n/\varepsilon^2)$ cells
 - Need to check intersection of n $T(a)$'s
 - $O(n^2/\varepsilon^2)$ time

Approximate Algorithm for $H()$

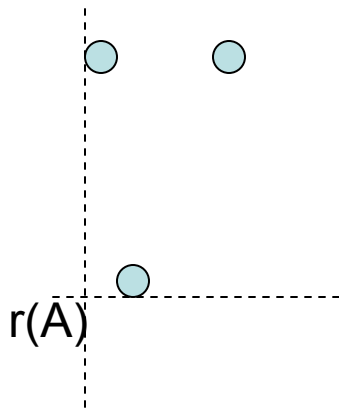
- Reference point: a point $r(A)$ such that if we set $t=r(B)-r(A)$, then

$$H(t(A), B) \leq c H_T(A,B)$$

- Note that t transforms $r(A)$ into $r(B)$
- This will give us a c -approximate algorithm with time $O(n \log n)$

Constructing a Ref Point

- $r(A) = (\min_{a \in A} a_x, \min_{a \in A} a_y)$



- Introduced in [Alt, Behrends, Blomer'91]
- Improved in [Aichholzer, Alt, Rote'94]

Theorem

Theorem: $r(A)$ is a reference point with $c=1+\sqrt{2}$

Proof:

- Consider optimal t , i.e., s.t. $H(t(A), B) = H_T(A, B)$
- What is $|\min_{a \in A} t(a)_x - \min_{b \in B} b_x|$?
- It is at most $H_T(A, B)$
- Same for $|\min_{a \in A} t(a)_y - \min_{b \in B} b_y|$
- Thus $\|r(t(A)) - r(B)\| \leq \sqrt{2} H_T(A, B)$
- We can translate $t(A)$ by a vector of length $\sqrt{2} H_T(A, B)$, so that $r(A)$ is aligned with $r(B)$

Exact matching under T+R

- Check if $DH_{T+R}(A,B)=0$
 - Can modify the algorithm to allow small error
 - Technique used in practice

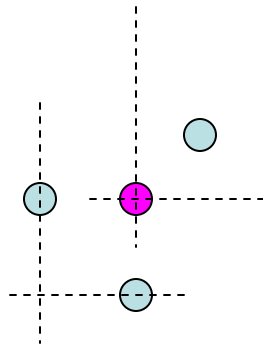
Algorithm

- Take any pair $a, a' \in A$, let $r = \|a - a'\|$
- Find all pairs $b, b' \in B$ such that $\|b - b'\| = r$
- For all such pairs
 - Compute t that transforms (a, a') into (b, b')
 - Check if $t(A) \subseteq B$

Analysis

- Choosing a, a' : constant time
- Enumerating b, b' : $O(n^2)$ time
- Checking match: $O(n P)$, where P is the number of pairs $b, b' \in B$ s.t. $\|b - b'\| = r$
- The bound for P is ...
- ...deferred to the next episode 😊

The $(1+\sqrt{2})$ Factor is Tight (for this reference point)



Face Recognition

