

Contour Correspondence via Ant Colony Optimization

Oliver van Kaick
GrUVi Lab

Ghassan Hamarneh
MIAL

Hao Zhang
GrUVi Lab

Paul Wighton
MCL

School of Computing Science
Simon Fraser University



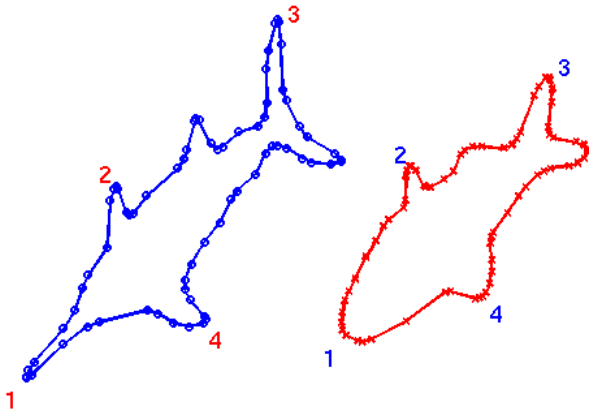
- 1 Introduction
- 2 Related work on correspondence
- 3 Review of the ACO framework
- 4 ACO for shape correspondence
- 5 Experimental results
- 6 Conclusions

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- Shape correspondence
 - Finding a meaningful matching between two shapes

Introduction

- Shape correspondence
 - Finding a meaningful matching between two shapes
- Focus: 2D contours



- Applications in
 - Computer graphics (shape analysis, morphing, and animation)
 - Computer vision (object tracking, recognition, and retrieval)
 - Medical computing (statistical shape modeling and analysis of anatomical structures)
- 3D shape matching and retrieval (Chen et al., 2003)

Solving contour correspondence

Common approach

- 1 Select feature points
- 2 Compute shape descriptors
- 3 Extract a matching

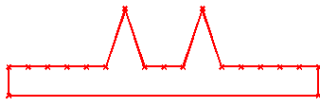
Common approach

- ① Select feature points
- ② Compute shape descriptors
- ③ Extract a matching
 - Greedy best matching
 - Bipartite matching
 - Iterative closest point (ICP) scheme
 - Dynamic programming under point ordering

Incorporating proximity information

- Most algorithms do not consider proximity information

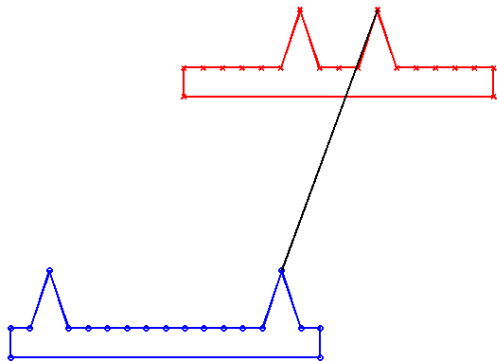
Proximity: if two points are close on one shape, their corresponding points in the second shape should also be close



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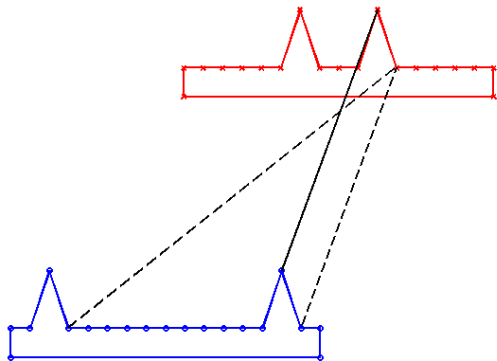
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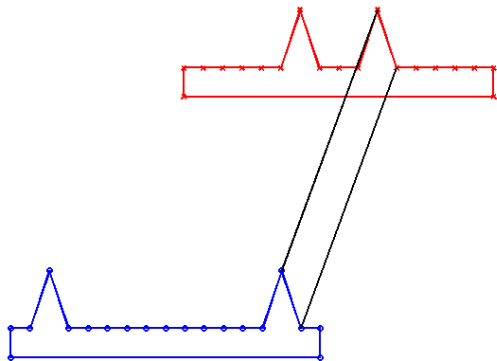
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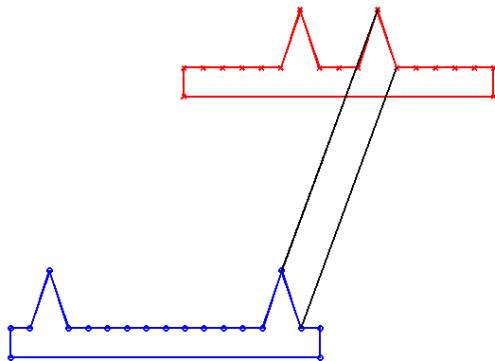
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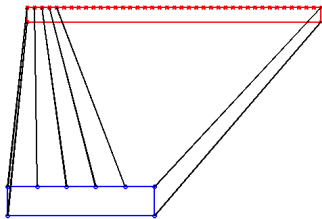


- Better handling of missing parts or a lack of salient features

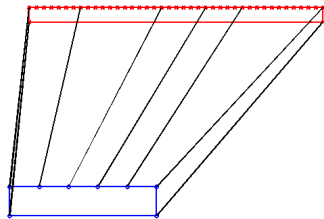
Incorporating proximity information

- Take advantage of the vertex ordering (contours)

Enforcing **order preservation** \neq



Enforcing **proximity**



- When **proximity** is incorporated, we can formulate point correspondence via the **Quadratic Assignment Problem (QAP)**
- **QAP** is one of the **most difficult** optimization problems
- **Ant Colony Optimization (ACO)** has had great success in solving this problem

- We formulate the general point correspondence problem in terms of **QAP** incorporating **proximity information**
- We propose the **first ACO algorithm** to compute the matching
- Applicable to **contours** and **unorganized 2D point sets**
- We extend the framework to enforce **order preservation** (contours)

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- Matching shape descriptors
 - 2D shape matching
 - Bipartite matching solved using the Hungarian algorithm
 - Integer constrained minimization (Maciel and Costeira, 2003)
 - Soft assign algorithm (Gold et al., 1998)
 - Preservation of binary neighborhood information (Zheng and Doermann, 2006)
 - QAP formulation (Berg et al., 2005)
 - Contour correspondence
 - Order preservation and dynamic programming (Liu et al., 2004, Scott and Nowak, 2006)

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- Without using local descriptors:
 - Physically-based approach (Sederberg and Greenwood, 1992)
 - Pattern matching in the Gaussian map (Tal and Elber, 1999)
 - Deformation-based edit distance (Sebastian et al., 2003)
 - Skeletal and shock graphs (Sundar et al., 2003, Siddiqi et al., 1999)
- Transform-based techniques (Shapiro and Brady, 1992, Sclaroff and Pentland, 1995, Bronstein et al., 2006, Jain et al., 2007)
- Group correspondence
 - Minimum Description Length (MDL) principle (Davies et al., 2002)

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- Routing, assignment, and scheduling
- Inspiration from nature
 - Individual ants have a simple behavior
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ACO facts

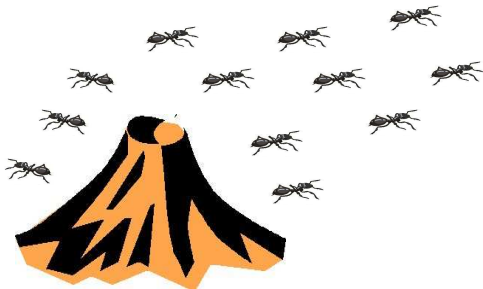
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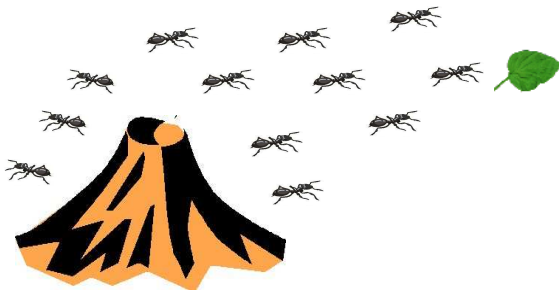


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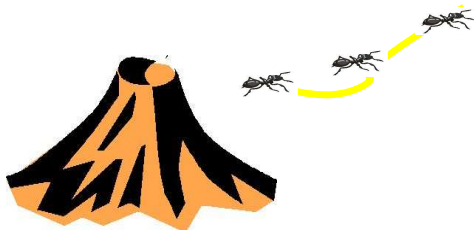


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Review of the ACO framework

- Problem modeled with a graph
- The solution search involves ants traversing this graph

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- ACO metaheuristic:

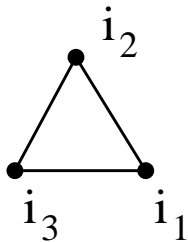
For each iteration:

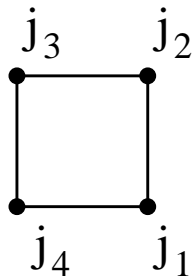
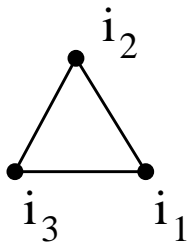
- 1 Traverse the graph (construct solution)
 - Heuristic information and pheromones
- 2 Evaluate solution
 - Objective function
- 3 Deposit pheromones on the edges of the graph
 - Quality of the solution

Advantages of ACO

- Probabilistic approach
- Heuristic information
- Escape from bad local minima
- Parallelizable

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Shapes to be matched

• i_2

• i_3 • i_1

• j_3 • j_2

• j_4 • j_1

i_1 ●

i_2 ●

i_3 ●

j_3 ● j_2 ●

● j_4 ● j_1

$i_1 \bullet$

$\bullet j_1$

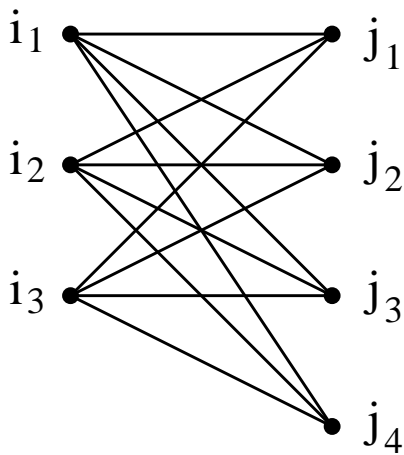
$i_2 \bullet$

$\bullet j_2$

$i_3 \bullet$

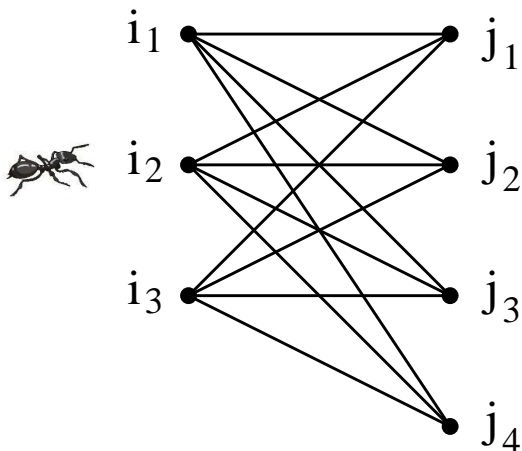
$\bullet j_3$

$\bullet j_4$



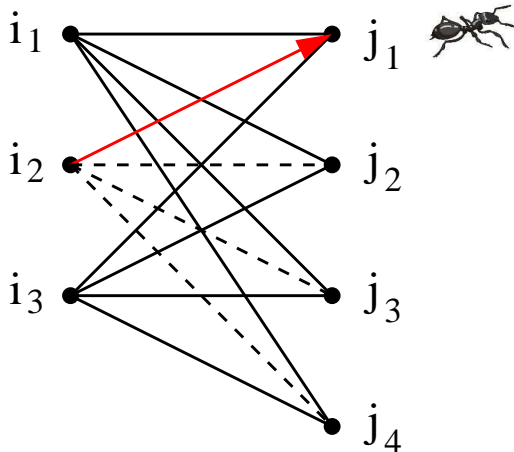
ACO graph

ACO for shape correspondence



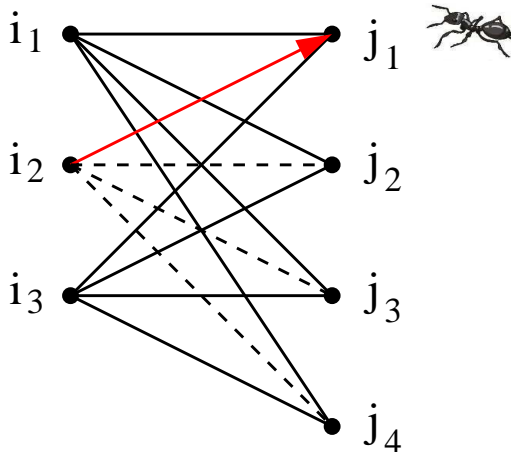
Iteration start

ACO for shape correspondence

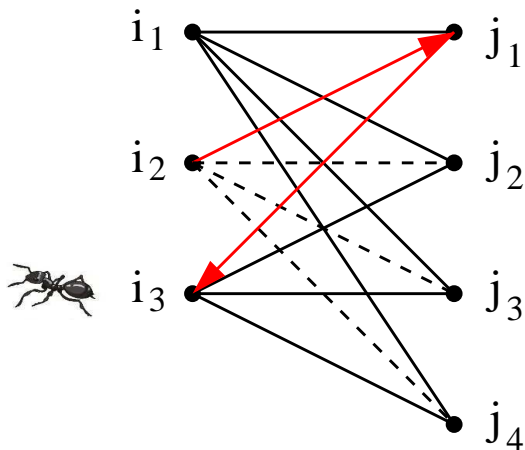


$$\pi(i_2) = j_1$$

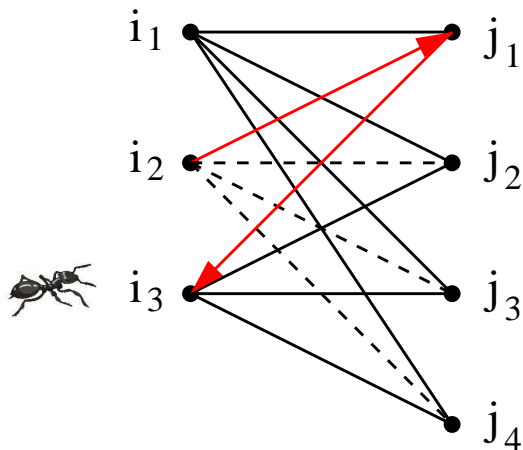
ACO for shape correspondence



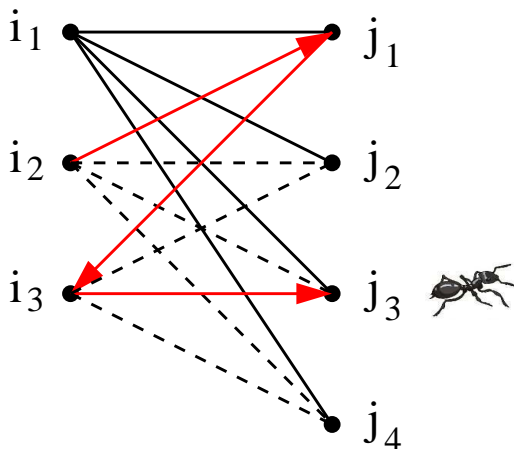
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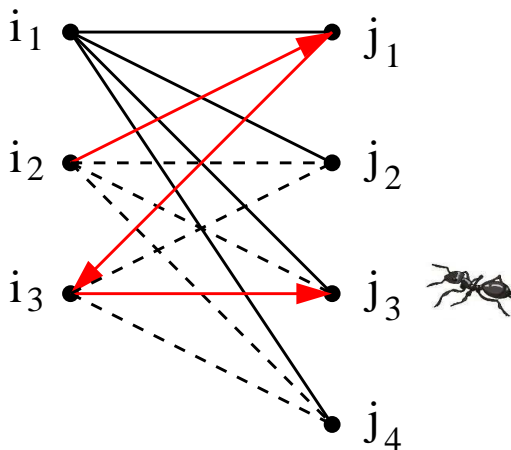


ACO for shape correspondence

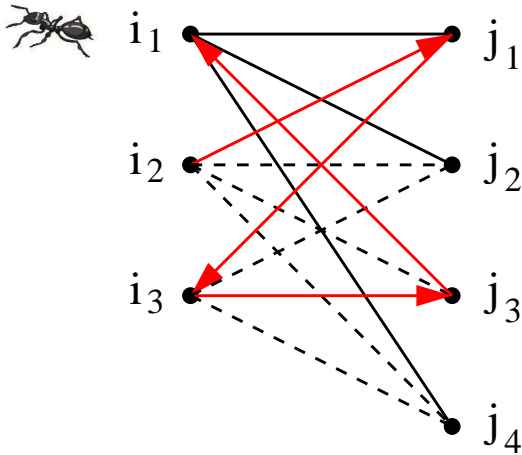


$$\pi(i_3) = j_3$$

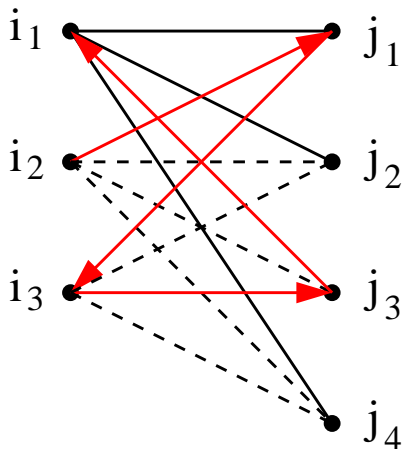
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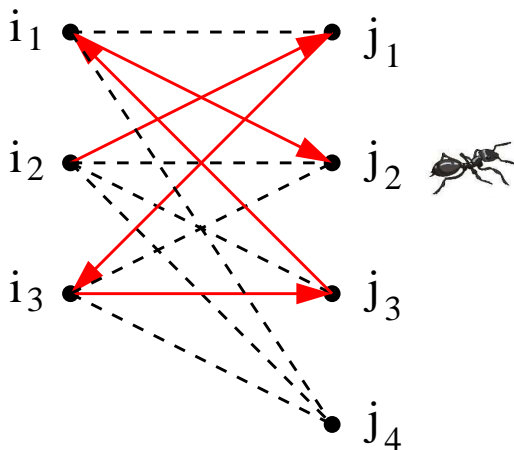
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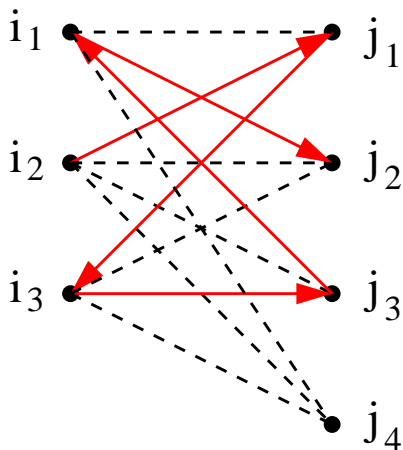


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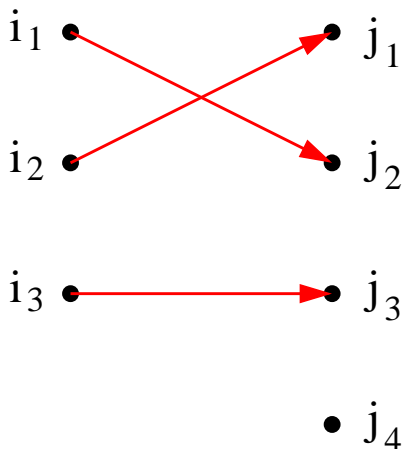
$$\pi(i_1) = j_2$$

ACO for shape correspondence



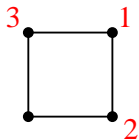
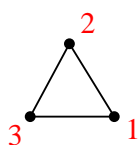
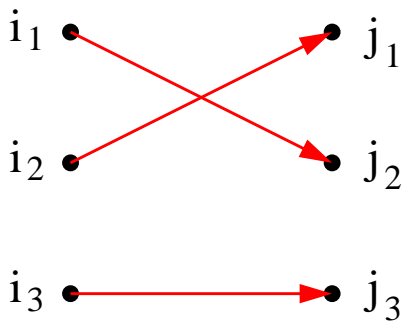
Iteration end

ACO for shape correspondence

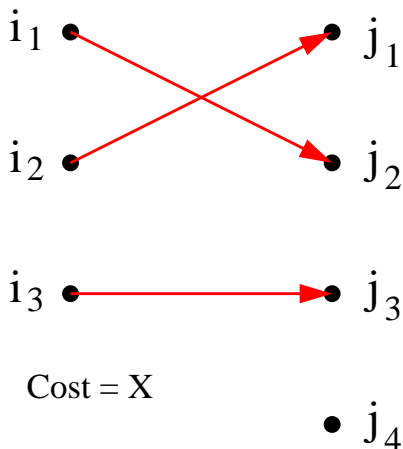


Correspondence obtained

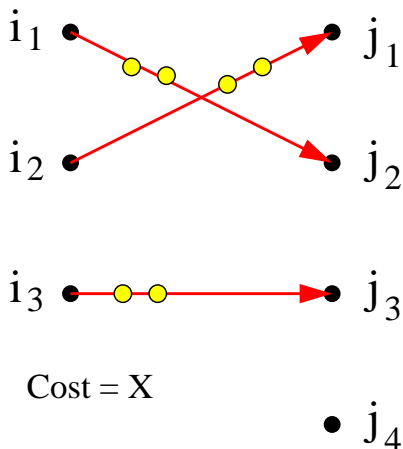
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- For a fixed number of iterations:
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Cost function: QAP formulation

$$\text{QAP} = \text{Descriptor distance} + \text{Proximity}$$

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$$\text{QAP} = (1 - v) \text{Descriptor distance} + v \text{Proximity}$$

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$$\text{Descriptor distance} = 1 - \frac{1}{|I|} \cdot \sum_{i \in I} \exp \left(- \frac{\text{dist}(R_i, R_{\pi(i)})^2}{\sigma_R^2} \right)$$

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$$\text{Proximity} = \sum_{i \in I} \sum_{i' \neq i \in I} |\text{dist}(i, i') - \text{dist}(\pi(i), \pi(i'))|$$

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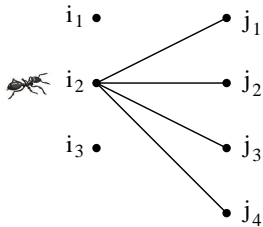
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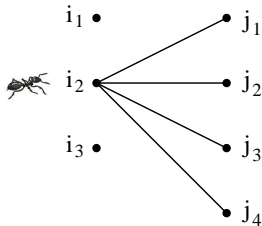
$$\text{Proximity} = \frac{\sum_{i \in I} \sum_{i' \neq i \in I} \exp \left(\frac{-\text{dist}(i, i')^2}{\sigma_I^2} \right) |\text{dist}(i, i') - \text{dist}(\pi(i), \pi(i'))|}{|I|(|I| - 1)/2}$$

Edge traversal



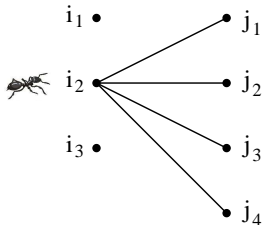
Edge probability:

$$p_{ij} = \alpha \text{Pheromones} + (1 - \alpha) \text{Heuristic}$$



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Heuristic information:

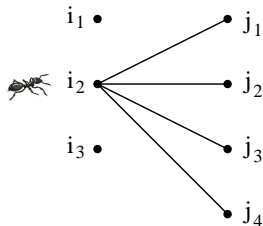
$$\text{Heuristic}^{-1} = \frac{\text{dist}(R_i, R_j)}{\text{Descriptor similarity}}$$

$$|\text{dist}(i, i') - \text{dist}(j, \pi(i'))| \times \text{Proximity to last vertex}$$

$$|\text{dist}(i, i'') - \text{dist}(j, \pi(i''))| \times \text{Proximity to second last vertex}$$

Edge probability:

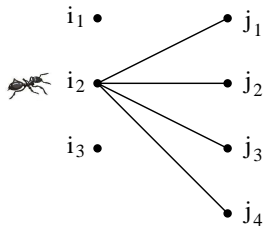
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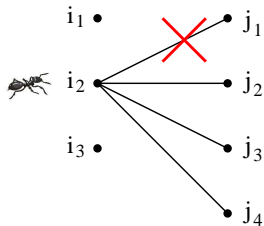
Heuristic information:

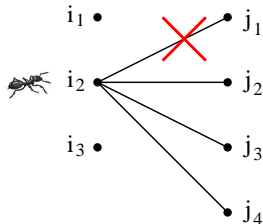
$$\text{Heuristic} = \left(e^{\frac{-\text{dist}(R_i, R_j)^2}{\sigma_R^2}} \right) \times \left(1 - e^{\frac{-\text{dist}(i, i')^2}{\sigma_I^2}} |\text{dist}(i, i') - \text{dist}(j, \pi(i'))| \right) \times \left(1 - e^{\frac{-\text{dist}(i, i'')^2}{\sigma_I^2}} |\text{dist}(i, i'') - \text{dist}(j, \pi(i''))| \right)$$

Order preservation



- Hard constraints: by removing edges that should not be traversed
 - Order preservation





- Hard constraints: by removing edges that should not be traversed
 - Order preservation
- Soft constraints: by modifying the probabilities

List of ACO parameters and values

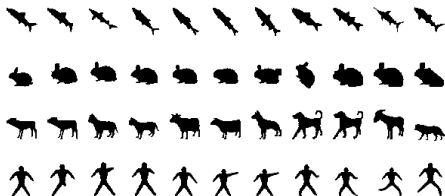
ACO Parameters	Symbol	Value
Number of ants	m	1
Number of iterations	T	1000
Influence of pheromones	α	0.3
Pheromone evaporation rate	ρ	0.1
Pheromone deposition constant	δ	0.01
Initial pheromone levels	τ_0	1
Minimum pheromone levels	τ_{\min}	$0.1 \cdot \frac{1}{ I }$
Influence of proximity	ν	0.7
Gaussian width in \mathcal{X}	σ_I	$0.1 \cdot l_{\max}$
Gaussian width in \mathcal{S}	σ_R	$0.1 \cdot R_{\max}$

Parameters used in our ACO algorithm and their chosen values

The values are fixed in all the experiments

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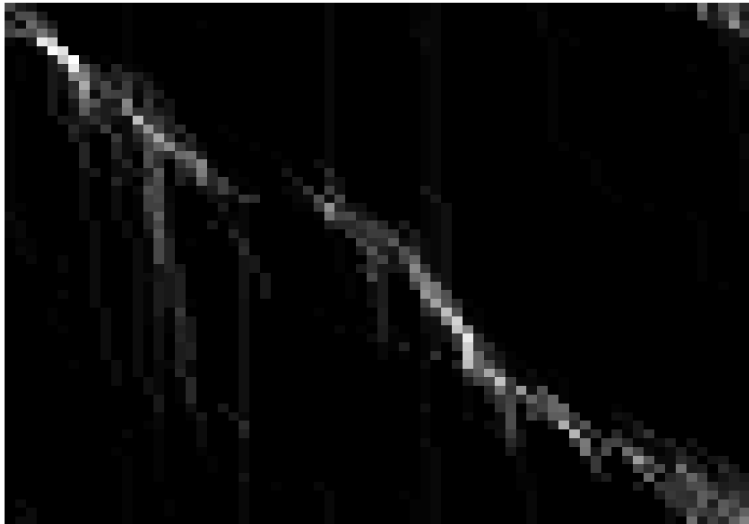
- Contours extracted from the Brown dataset (Sharvit et al., 1998)



- Shape context (rotation-variant) descriptor (Belongie et al., 2002)

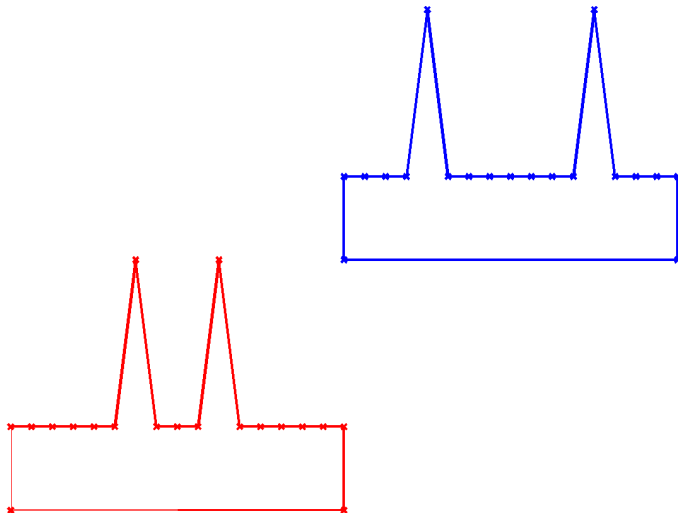
ACO pheromone deposition

Iteration 300 - Cost of best matching so far: 0.25704

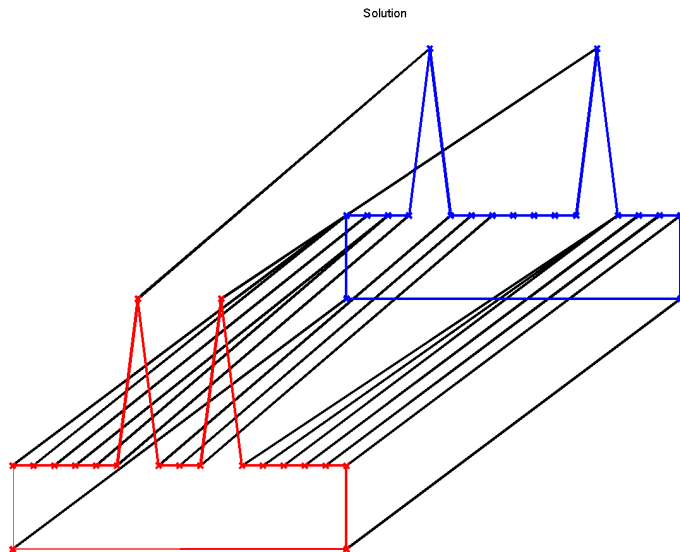


ACO solution computation

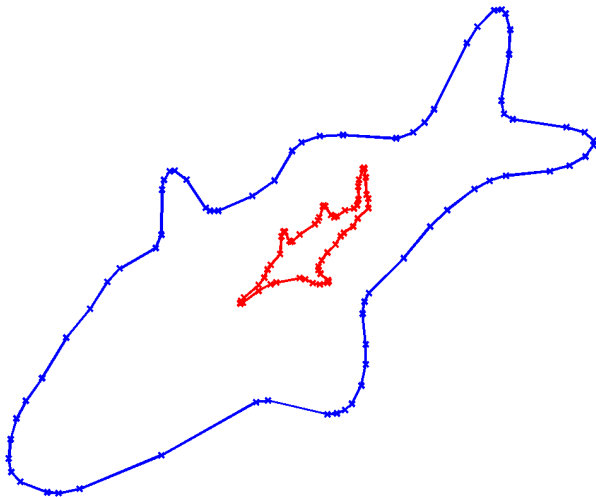
Shapes to be matched



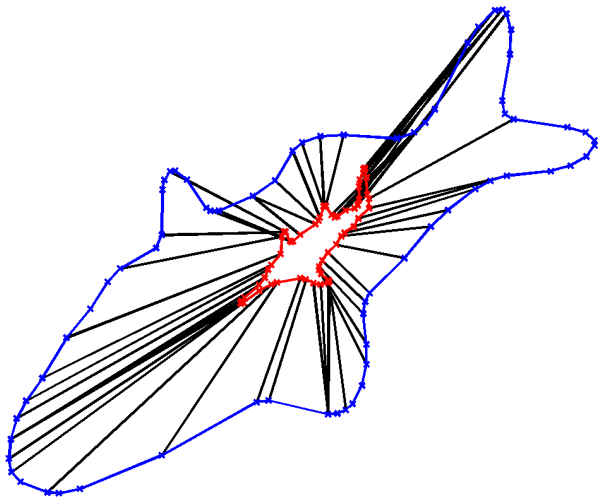
ACO solution computation



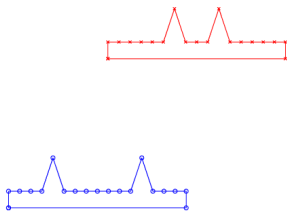
Shapes to be matched



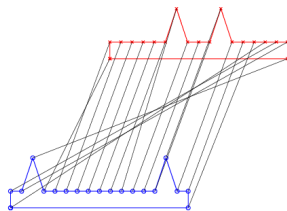
Solution



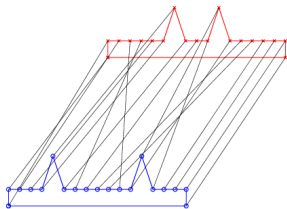
Order preservation (OP) vs. proximity



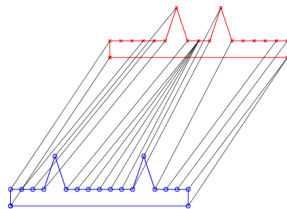
(a) Contours to match



(b) OP

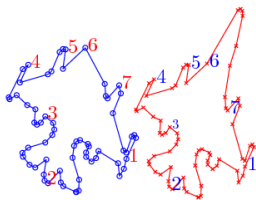


(c) ACO (Proximity only)

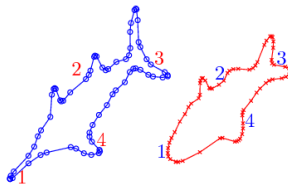


(d) ACO (Proximity + OP)

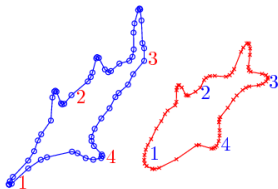
Handling of occlusion or missing parts



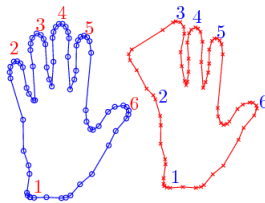
(a)



(b)



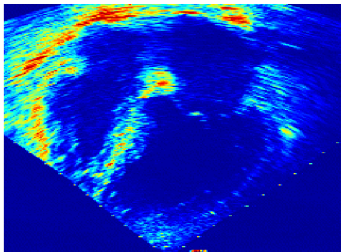
(c)



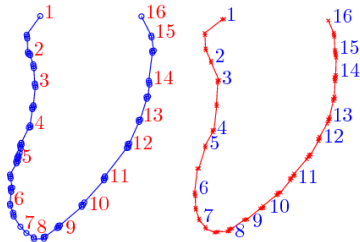
(d)

Matchings computed by ACO for contours
with occlusion or structure change

Handling of open contours



(a) Frame



(b) Matching

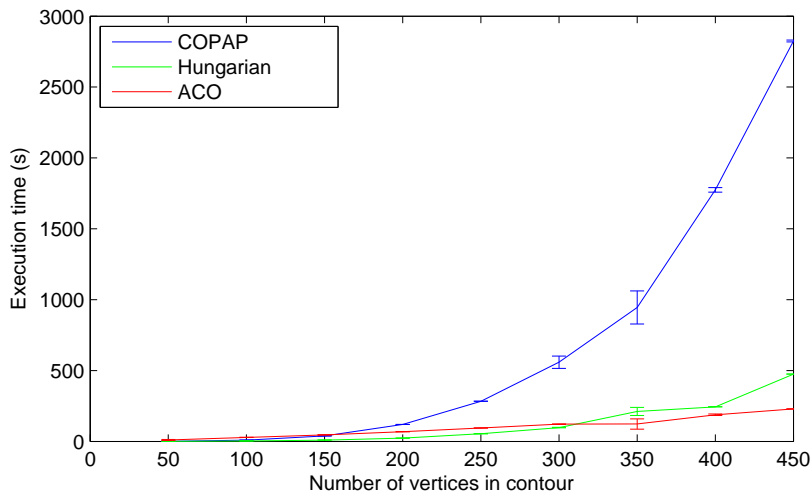
Matching computed by ACO for an open contour of a left ventricle

Evaluation against ground-truth correspondence

- Ground truth provided by a human user
- Error is the sum of geodesic distances between corresponded vertices and ground truth (Karlsson and Ericsson, 2006)
- Compared to Hungarian and COPAP (Scott and Nowak, 2006)

Shape class	Hungarian	COPAP	ACO
Airplanes	223.16	32.55	13.02
Fish	56.85	21.67	22.80
Four-legged	235.57	32.58	25.48
Hands	375.94	94.86	121.95
Humans	482.27	53.75	20.95
Rabbits	190.01	80.01	53.44
Stingrays	30.55	5.88	5.16
Tools	204.36	35.29	22.48

Deviation from ground truth



Execution time comparison between Hungarian, COPAP, and ACO

- 1 Introduction
- 2 Related work on correspondence
- 3 Review of the ACO framework
- 4 ACO for shape correspondence
- 5 Experimental results
- 6 Conclusions**

Conclusions and future work

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- The QAP is solved using the proposed ACO algorithm

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- Future work
 - Extension to 2D manifolds
 - Parallelization

Thank you!

Funding for this project was provided by

- Faculty of Applied Sciences
Simon Fraser University



gruvi  graphics + usability + visualization