

# GRAPP 2020 - 15<sup>th</sup> International Conference on Computer Graphics Theory and Applications

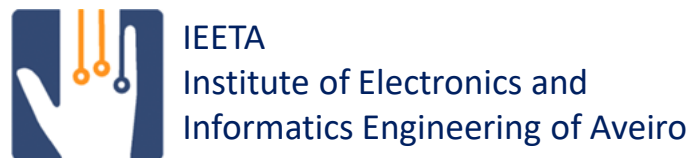
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## Matching-Aware Shape Simplification

### Authors

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- **Spatiotemporal Data**

- Moving Objects
  - Moving Points
  - Moving Lines
  - Moving Regions

- **Sample uses**



Satellite images of the evolution of an iceberg



Evolution of burned areas

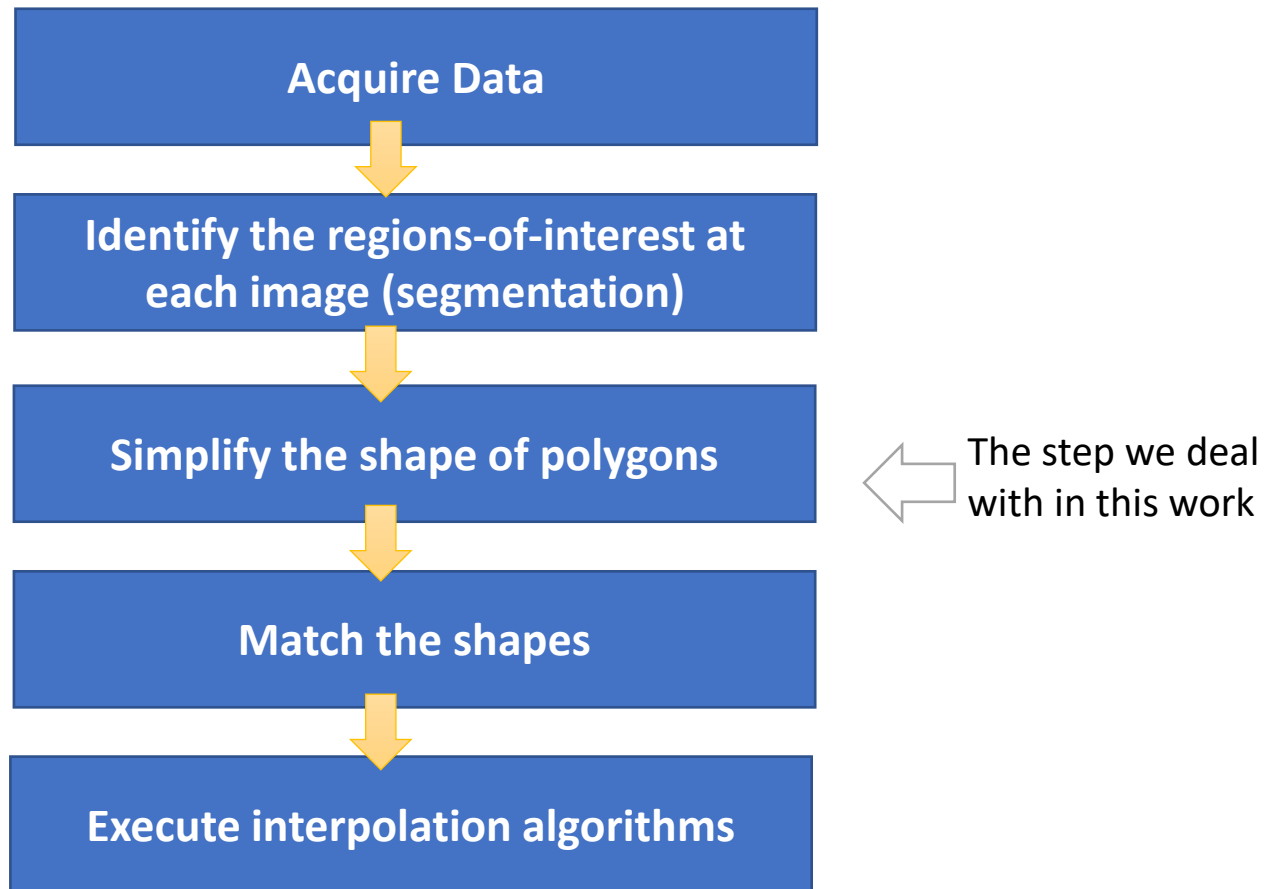
## • Continuous Representation

- Sliced representation (Forlizzi, L. et al. 2000) - *moving region* is represented by:  
(i) geometry + (ii) function (to represent the object's evolution)
- *Region interpolation problem*: With object representations at two timestamps, use a *function* to interpolate the representations



Simulating the burnt area evolution  
using a region interpolation algorithm

- **Data Preparation Workflow**



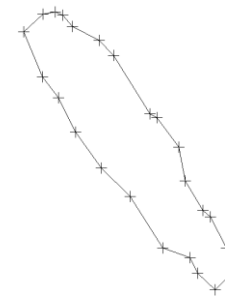
- In simplification algorithms focused in one-by-one simplification...
- May select vertices that are distant from their corresponding in another polygon
- Corresponding shapes may have different number of vertices (but some interpolation algorithms require the same number of vertices)



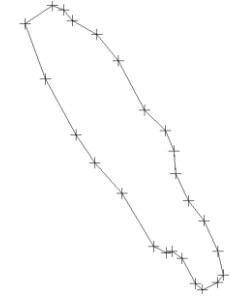
(a) Raw data - ice01



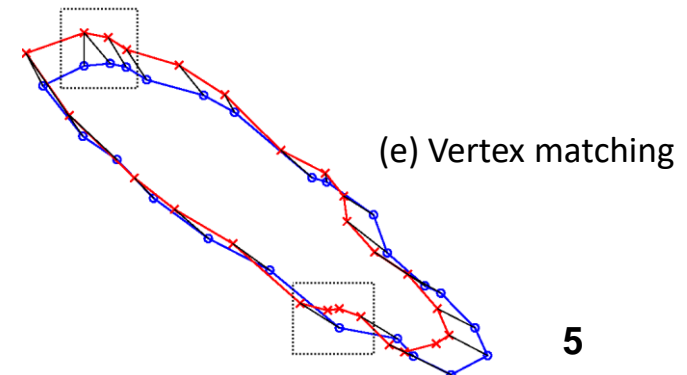
(b) Raw data - ice02



(c) Simplified Polygon  
- ice01



(d) Simplified Polygon  
- ice02




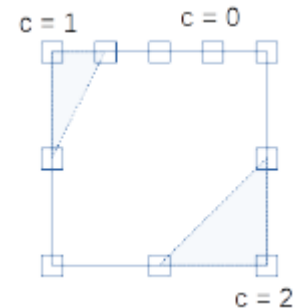
(e) Vertex matching

- Simplify two shapes simultaneously
- Considers the importance of vertices
  - to each shape representation
  - to inter-shape correspondence
- Objectives
  - Reduce adjustments during matching (e.g. adding extra vertices)
  - Provide locally-aware vertices to matching algorithms (to reduce vertex matching complexity)

- Considering two geometries  $P$  and  $Q$ , and the vertices  $p$  and  $q$  ( $p \in P, q \in Q$ )
- For each vertex  $p$ , the loss on removing  $p$  is:

$$\mathit{cost}_p = \max(\mathit{cost\_single}_p, \mathit{cost\_matched}_p * \mathit{t\_factor})$$

- $\mathit{cost\_single}_p$  = cost to remove for a single shape (e.g. the area of the triangle given by  $p, p-1$  and  $p+1$ ). 



- $\mathit{t\_factor}$  = preference to local or temporal information
- $\mathit{cost\_matched}_p$  = cost for loss of feature representation on  $Q$ :

$$\mathit{cost\_matched}_p = \max(\mathit{cost\_unique\_feature}_p, \mathit{cost\_matched\_feature}_q)$$

A significant vertex represents either:

- A feature present in Q and not in P

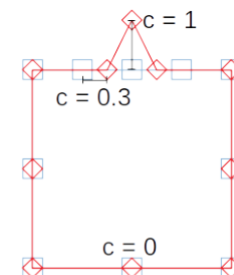
$$\mathit{cost\_unique\_feature}_p = \min(d_{pq})$$

- A vertex on P needed to morph into the feature of Q:

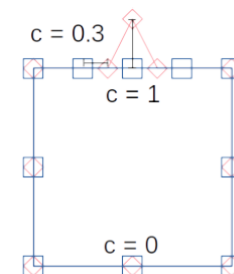
$$\mathit{cost\_matched\_feature}_p = \max(d_{pq})$$

$$d_{pq} = \min(d_{kq})$$

- Our algorithm starts removing the vertex in  $P$  with the lowest cost, then remove the one from  $Q$ , then remove another from  $P$ , and so on until  $P$  and  $Q$  have a desired size.



(a) Distinct features cost on Q (P on background)

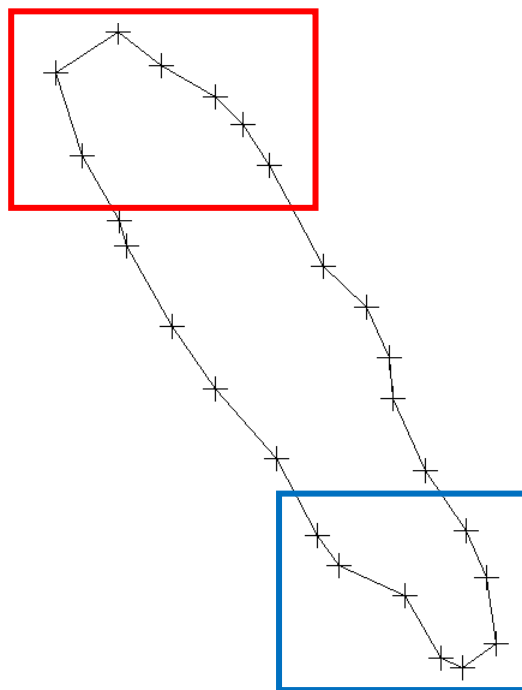
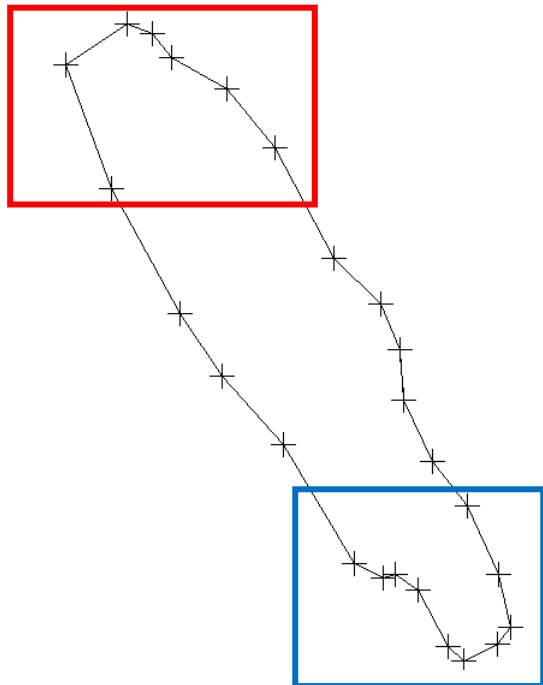


(b) Matching vertices on P (Q on background)

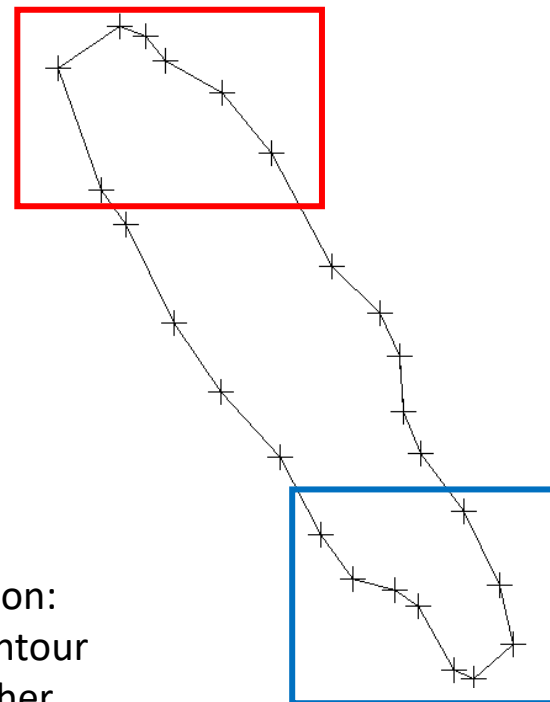


- **Simplification – Real World Data (Iceberg B-15a)**

Douglas–Peucker algorithm



Visvalingam-Whyatt algorithm

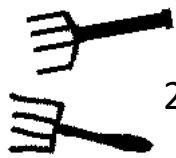


Matching-Aware Simplification:  
density of points along the contour  
more similar than that of other  
methods

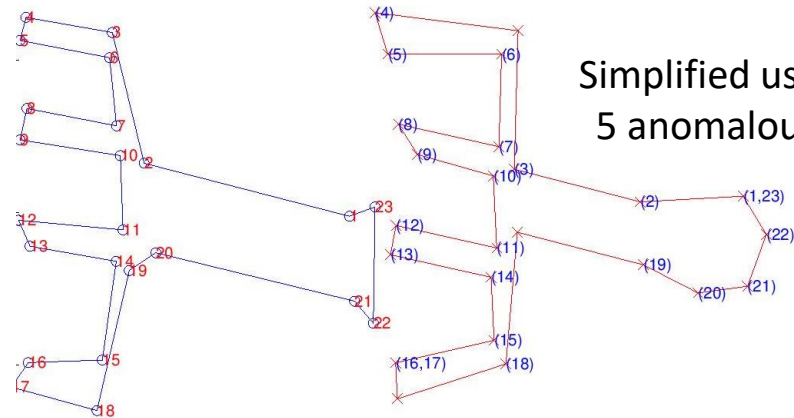
- **Simplification and Matching (216 Binary Shape Database)**
  - Dataset: 216 images from 18 classes
  - Executed each of the evaluated simplification algorithms
  - Executed the matching using the algorithm from Van Kaick et al.\*
  - Recorded the number of *anomalous* points during matching:
    - vertices without correspondence
    - vertices with multiple correspondences

\* O. van Kaick, G. Hamarneh, H. Zhang and P. Wighton, "Contour Correspondence via Ant Colony Optimization," *15th Pacific Conference on Computer Graphics and Applications (PG'07)*, 2007, pp. 271-280.

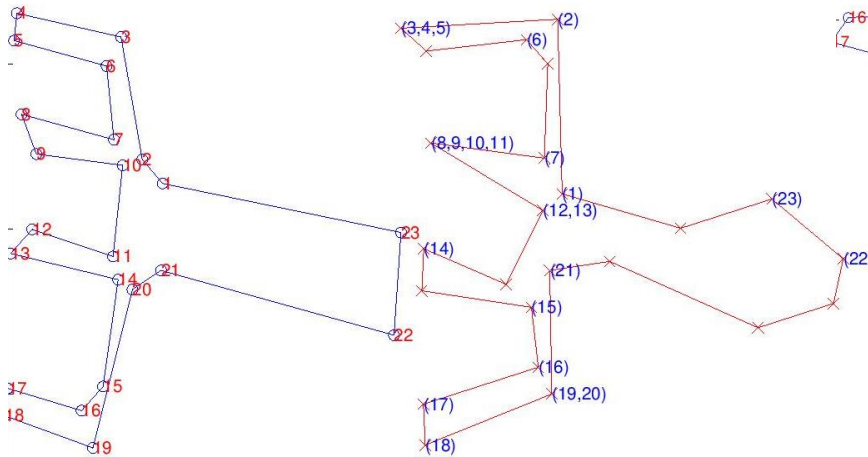
## • Simplification and Matching (216 Binary Shape Database)



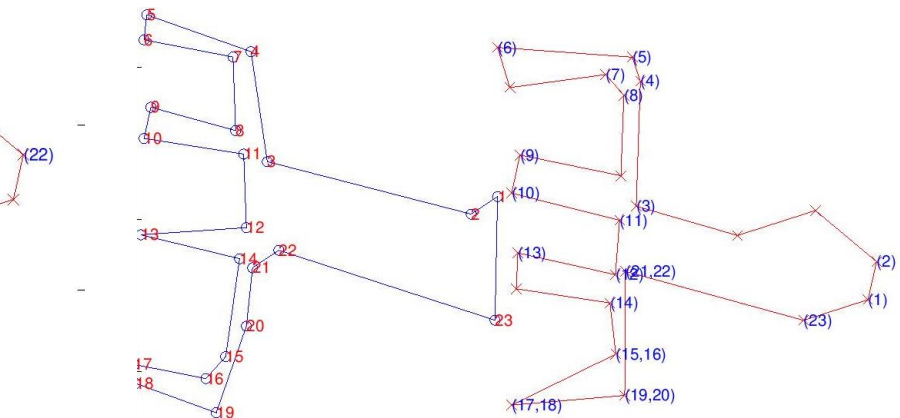
38 points in the first fork  
23 points in the second fork



Simplified using MAS:  
5 anomalous points

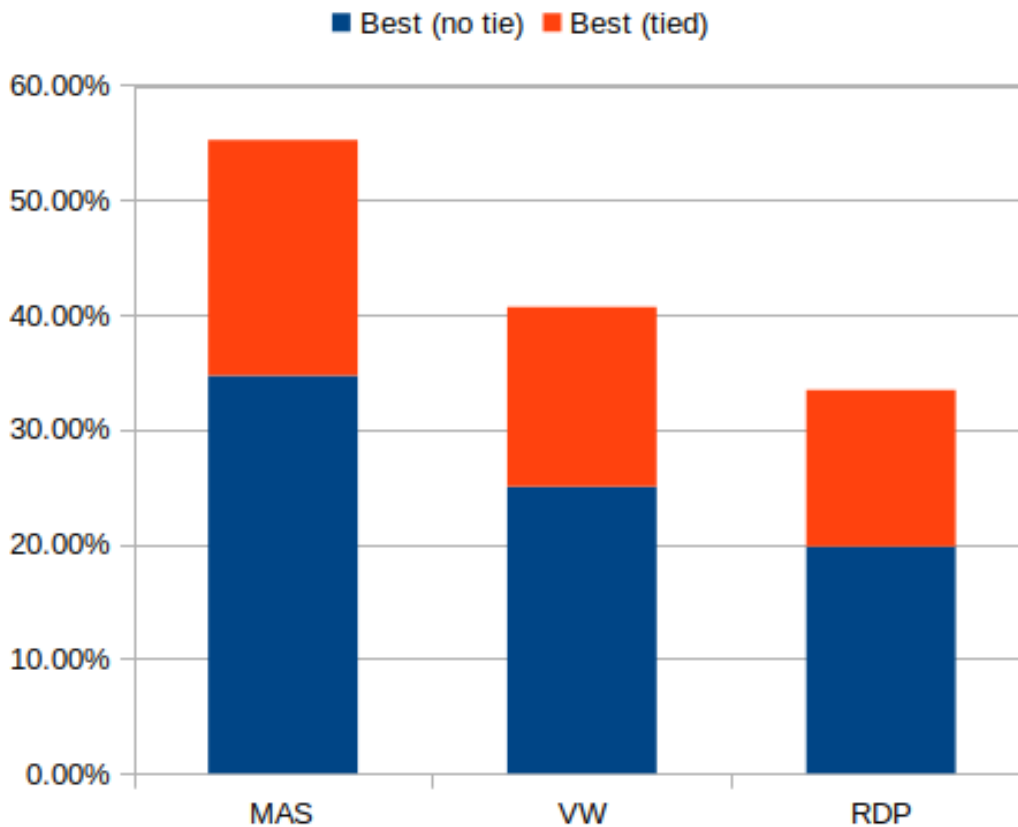


Simplified using Visvalingam-Whyatt algorithm:  
12 anomalous points



Simplified using Douglas-Peucker algorithm:  
9 anomalous points

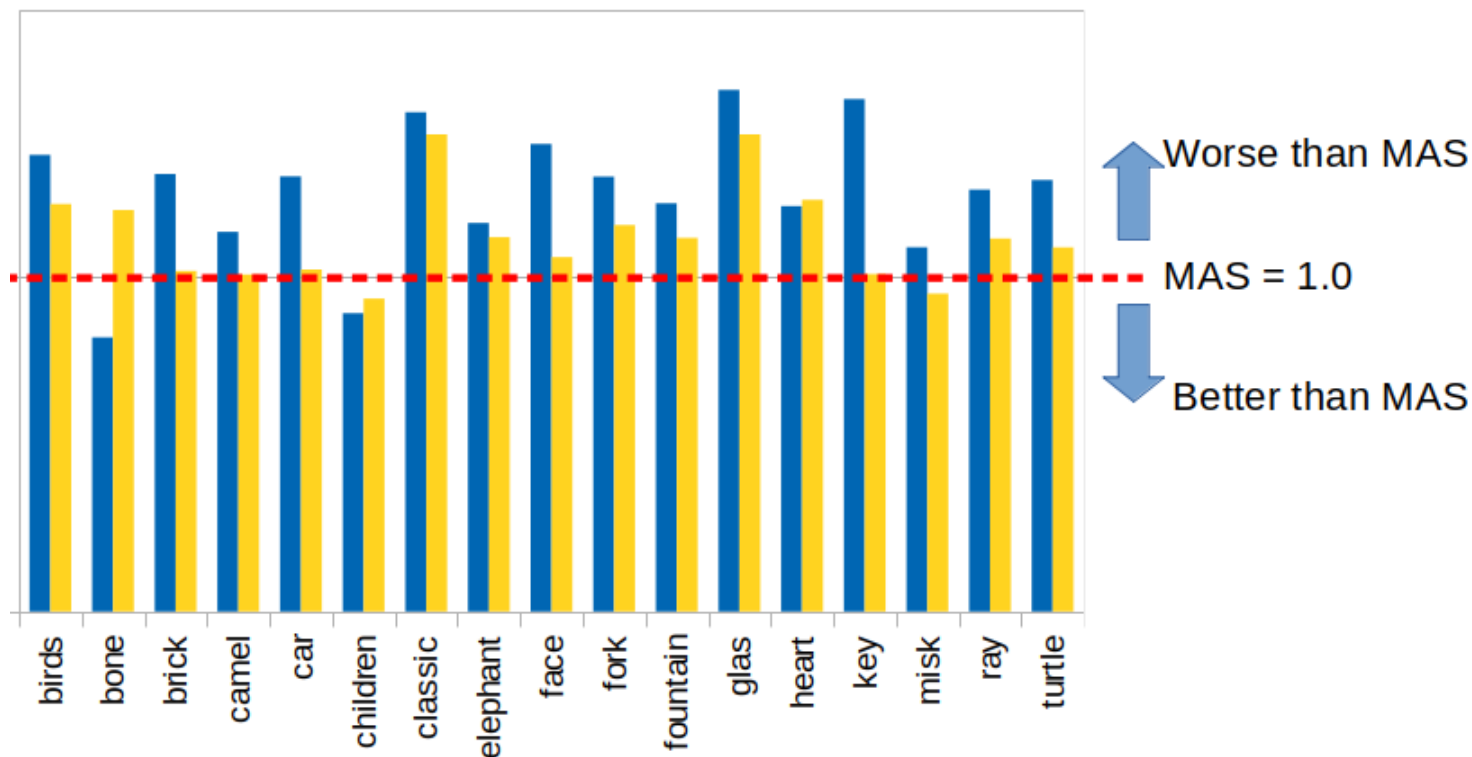
- **Simplification and Matching (216 Binary Shape Database)**



*Best* is the one that leads to a matching with the lowest number of anomalous points

- **Simplification and Matching (216 Binary Shape Database)**

Gap of performance per class (number of *anomalous* points):  
MAS has small defeats and big victories



- Matching-Aware Simplification
  - reduces the number of anomalous points during matching
  - can be combined with any matching algorithm
- Future work
  - Study the combined execution of simplification and matching
  - Study the simplification/matching in a wide sequence of polygons (not just two) to generate more natural interpolations
- New datasets (case studies) for moving regions are welcome!

**Thank you!**

**Questions?**

Project homepage: <http://most.web.ua.pt/>