SIFT descriptor to set landmark on biological images

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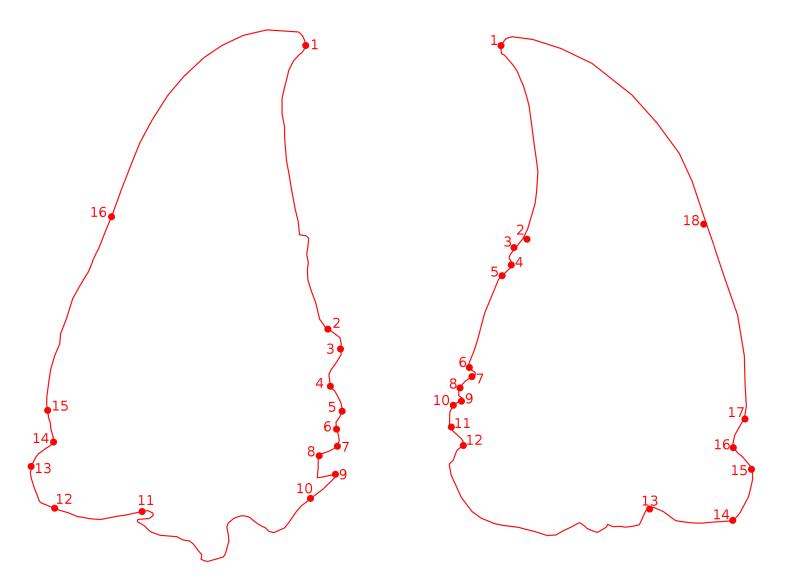


Context

- Morphometry analysis is a way to characterize the shape variations of the organisms,
- Morphometric characteristics have been used to evaluate the evolution of an organism, by finding new or sharpening the definition of old,
- Morphometrics are also used to classify the objects to different groups in a family.

Manual landmarks

- Morphometric landmarks are points that are a kind of points of interest,
- Landmarks are along an image outline and contain a lot of important information,
- They are defined by the biologists.



How to locate the landmarks automatically?

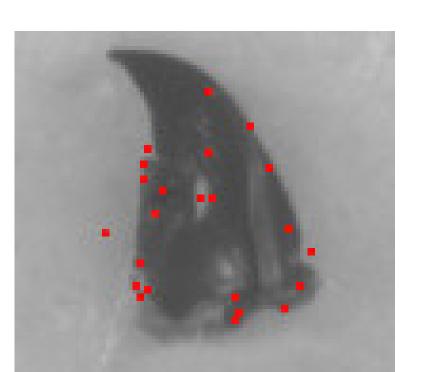
SIF1

SIFT[4] is used to extract distinctive features from the images. It includes four steps:

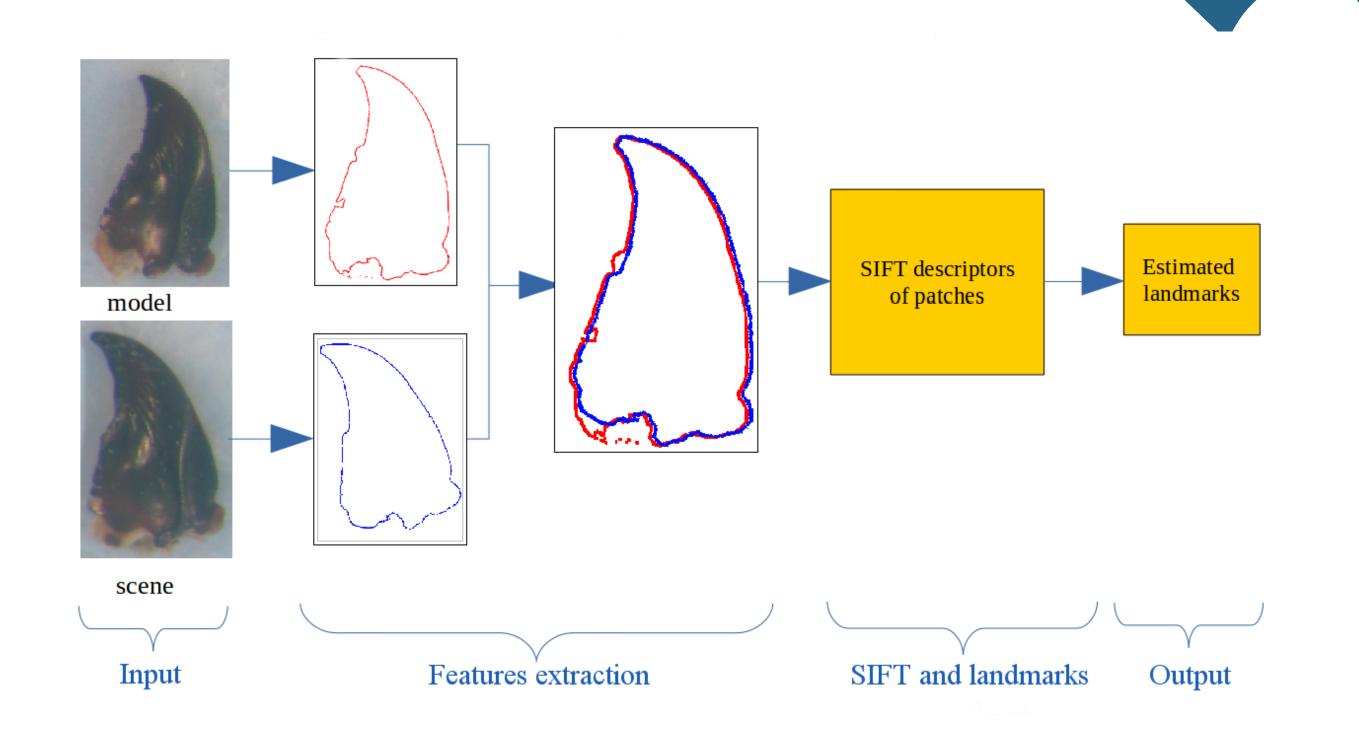
- Scale-space extrema detection
- Keypoints localization
- Orientation assigment
- Keypoint descriptor

The original SIFT outputs many candidates for landmarks.

<u>Solution:</u> Limiting the searching space before computing the SIFT descriptors.



Proposed method



Segmentation

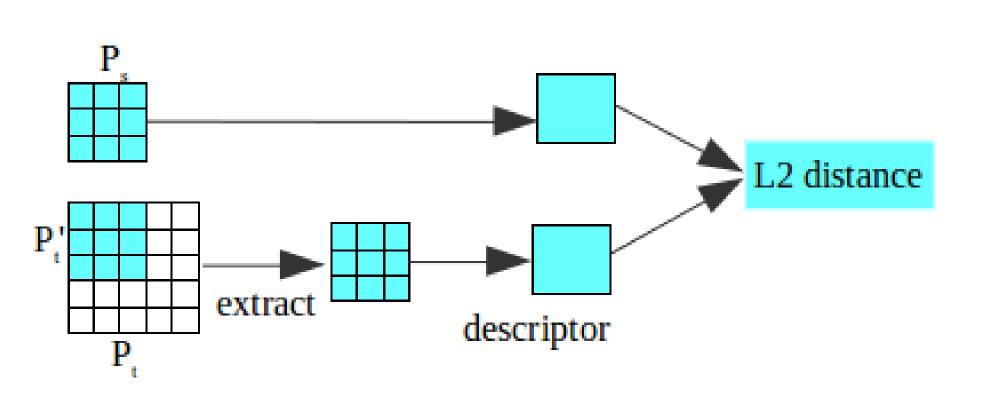
- Converting the image to binary by applying binary threshold. The threshold value is determined by analysing histogram[3].
- Contours points are extracted by Canny algorithm[1].
- The threshold ratio in Canny: $T_{lower} = (1/3) \times T_{upper}$

Registration

Two lists of contour points from segmentation step are registered by applying Principal Component Analysis[2, 5] Iteration (PCAI).

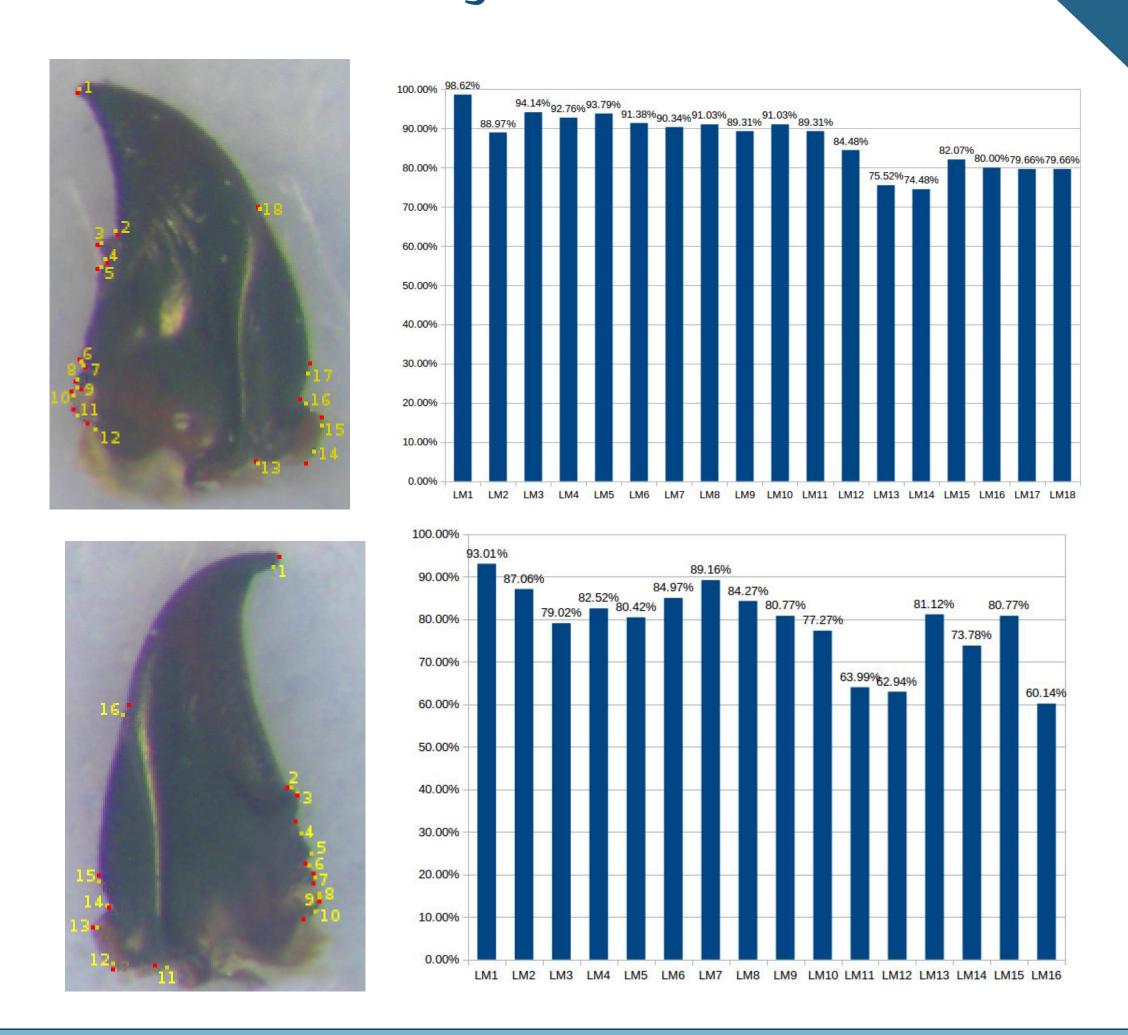
- 1. Compute the centroid point and principal axis of contours,
- 2. Compute the transformation values between two lists of contour points,
- 3. Register two images,
- 4. Sort the contour points followed y-direction,
- 5. Select a subset of contour points and repeat step 1,
- 6. PCAI stop automatically when the angle difference between two lists of contour points is less than 1.5 degree.

SIFT and landmarks



- 1. A patch P_m is initialized at each manual landmark of source image (size of 9×9),
- 2. Calculate the SIFT descriptor for P_m ,
- 3. At the same position in target image, a patch P_s is created (size of 36×36),
- 4. For each pixel in P_s , a patch P_s' is extracted with the same size of P_m ,
- 5. Calculate the SIFT descriptor for P'_s ,
- 6. Compute the distance between the descriptor of P_m and each P_m' ,
- 7. The process stops when all the pixels in P_s are considered. Keep the pixel that has the minimum distance with P_m .

Results on left and right mandibles



Bibliography

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