



SIFT descriptor to set landmarks on biological images

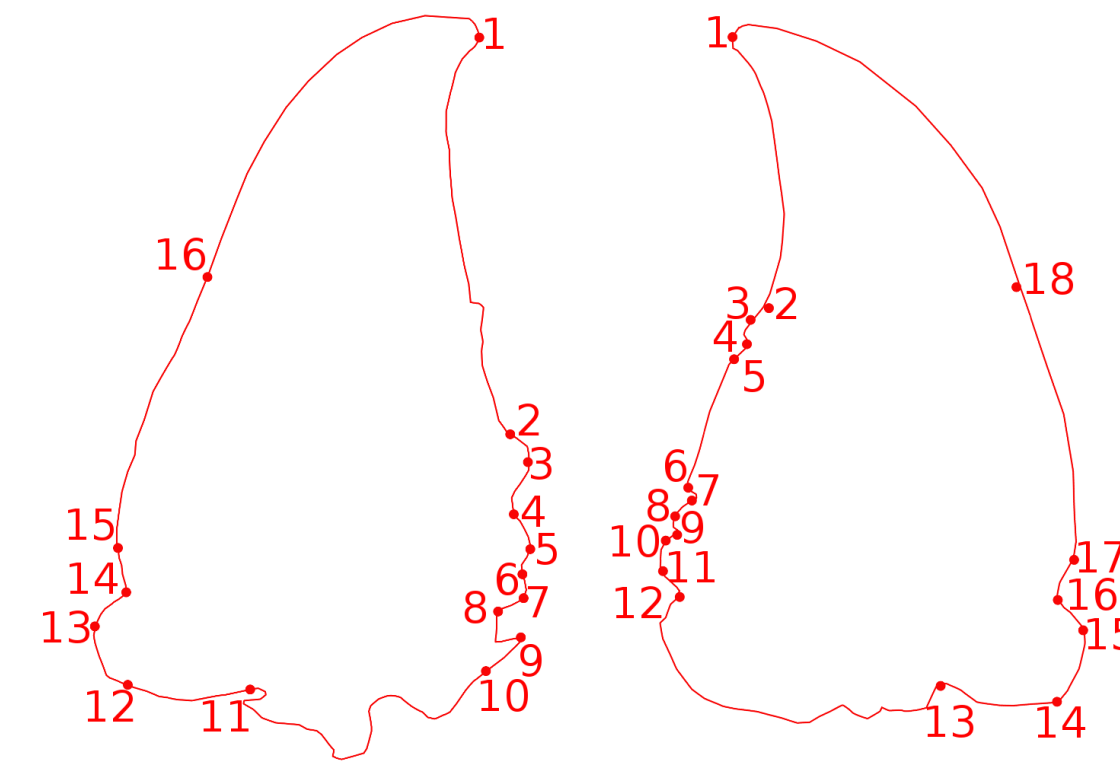
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Context

Morphometry analysis is a way to distinguish the characteristics of organisms, i.e. *shape, size, form,...*. It is used to appreciate the covariances between the ecological factors and the organisms. Landmark-based morphometry is known as one of the approaches to analyze the characteristics of organisms. Finding enough landmarks can give to biologists a comprehensive description of the organism. This work focuses on the automatic identification of landmarks on 2D biological images.

Landmarks

- Morphometric landmarks are points of interest in the biological object. They usually stay along the outline of the image.
- Landmarks characterize specificities through the shape most often linked to biological information,
- They are usually **defined manually** by biologists,
- Images at the right side show manual landmarks in **beetle mandibles** belonging to our sample:
16 and 18 manual landmarks have been defined for each left mandible and right mandible, respectively.



How to locate the landmarks automatically?

Mandibles and manual landmarks

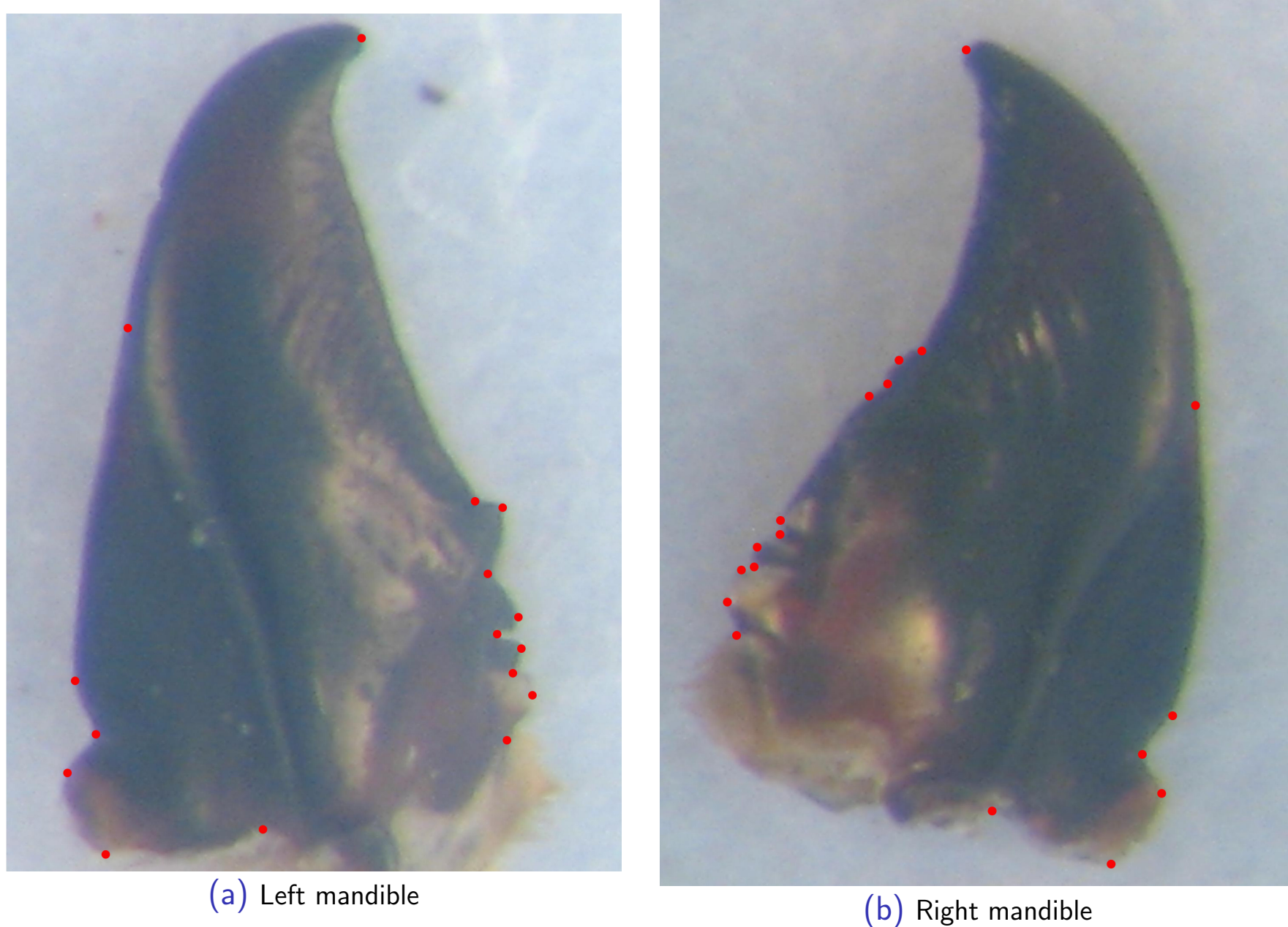
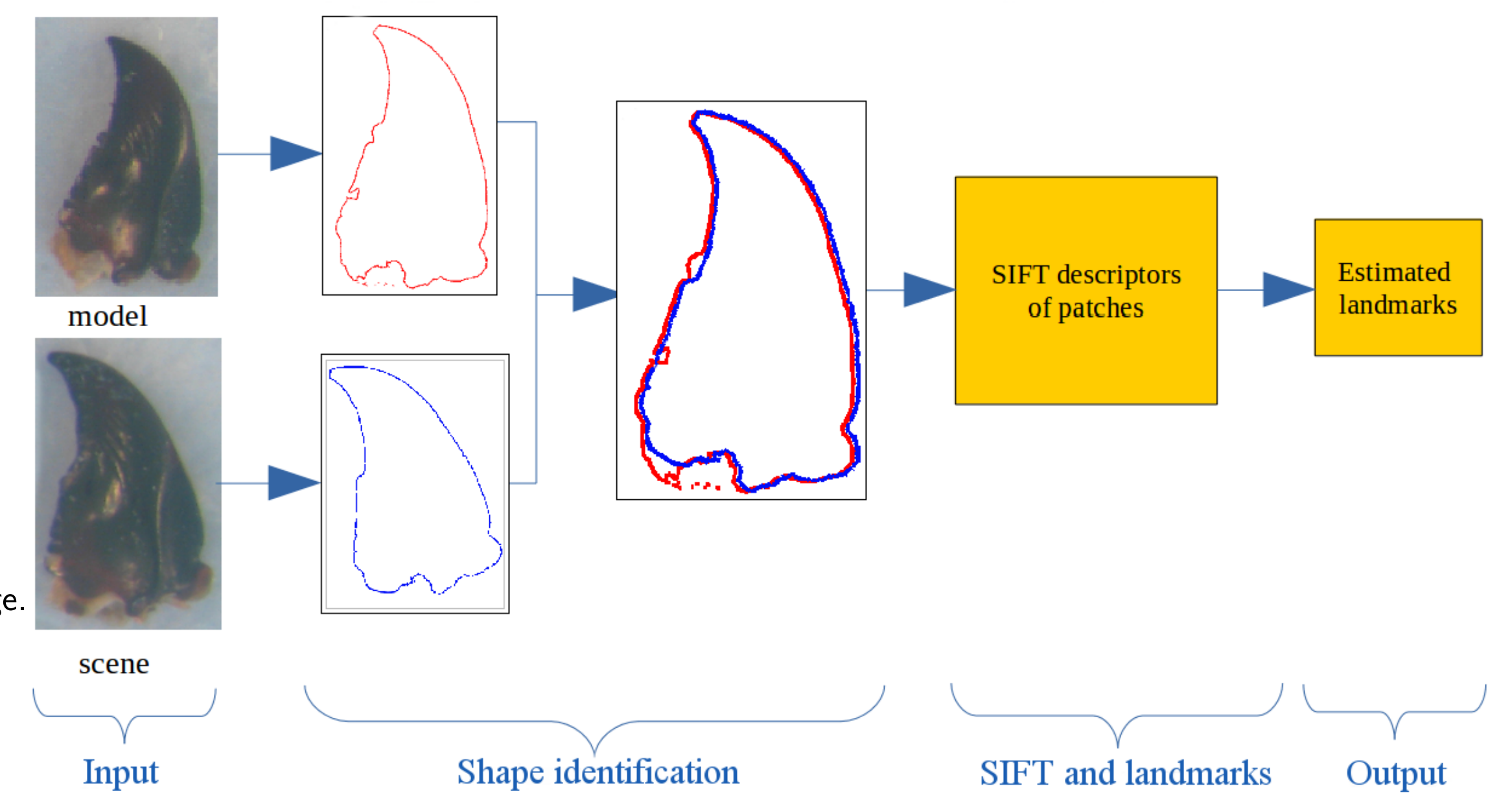


Figure: Example of beetle mandibles from the studied data set with manual landmarks.

Proposed method

- **Input:**
 - A model image
 - The manual landmarks of model image
 - A scene image
- **Output:**
 - Landmarks of scene image
- **Steps:**
 - Shape identification: segmentation and registration
 - SIFT and landmarks: Extract the patches, calculate the SIFT descriptors and estimate the coordinates of landmarks on the scene image.



Segmentation

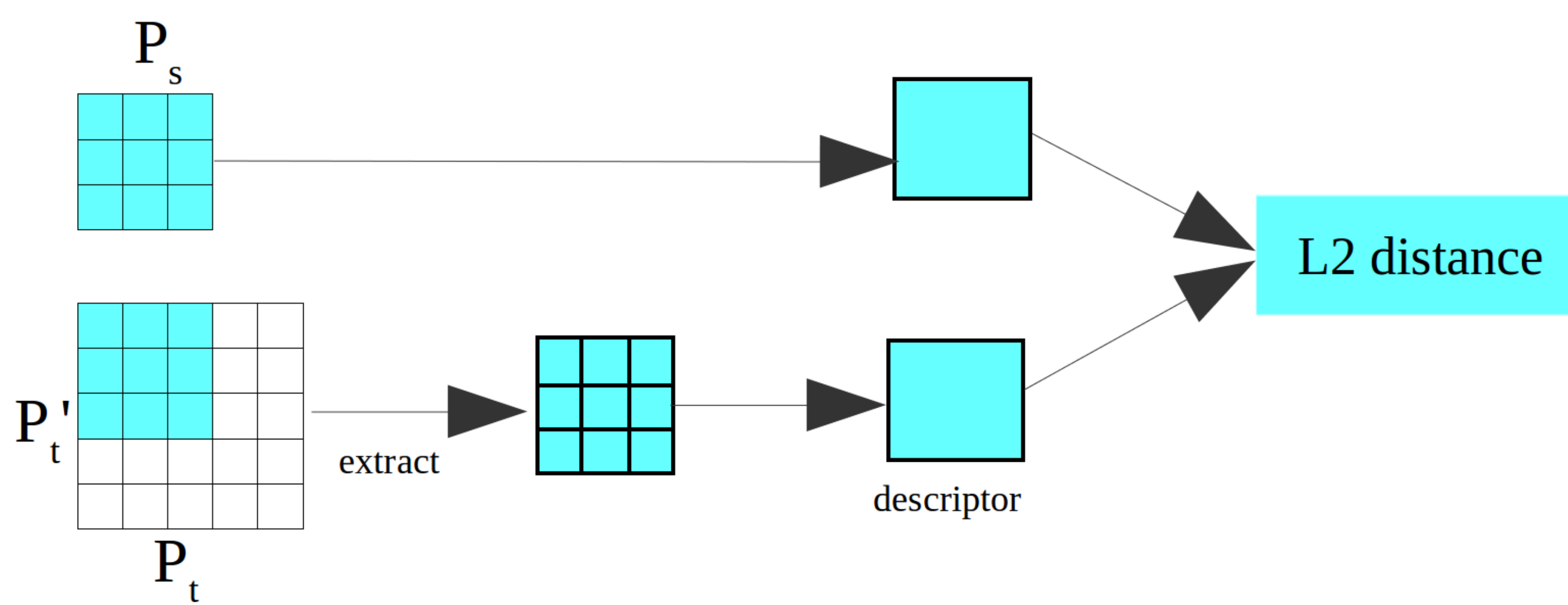
1. Converting the image to binary one by applying a threshold determined by histogram analysis [1],
2. Contours points are extracted by Canny algorithm [2]. The thresholds ratio in Canny: $T_{lower} = (1/3) \times T_{upper}$, in which T_{lower} equals to the threshold value in step 1.

Registration

Model and scene images are segmented to extract the contours points. The contours points are registered by applying **Principal Component Analysis [3] Iteration (PCAI)**.

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

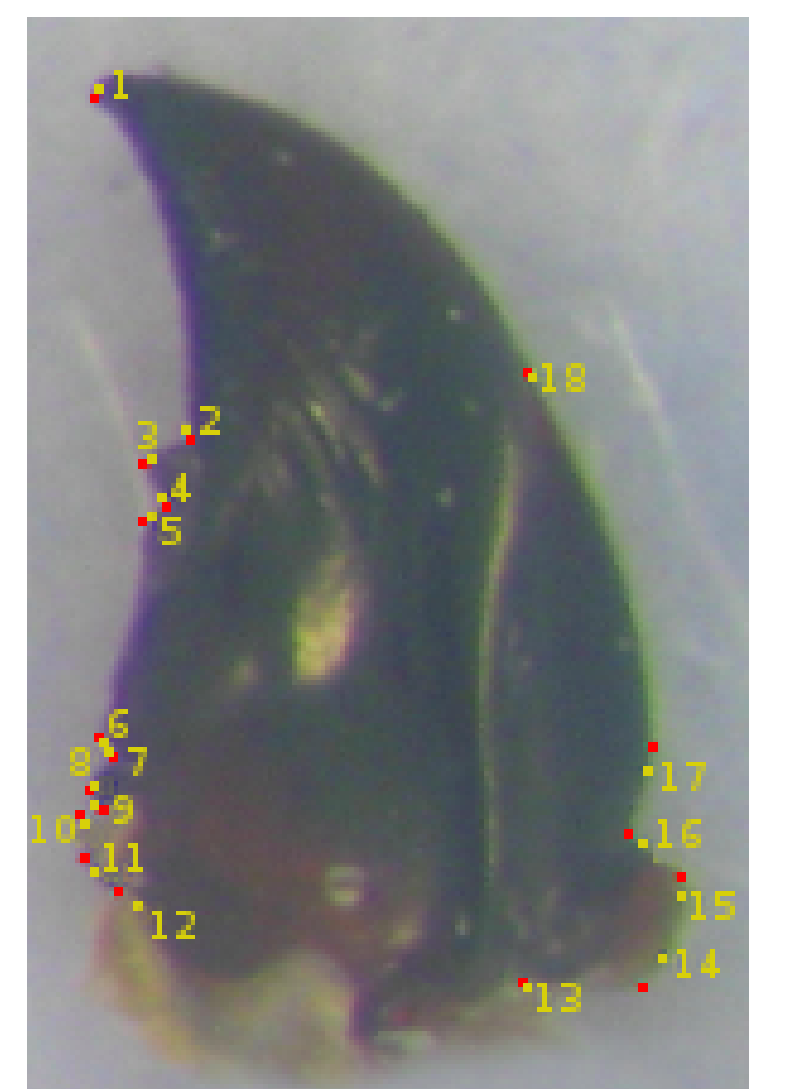
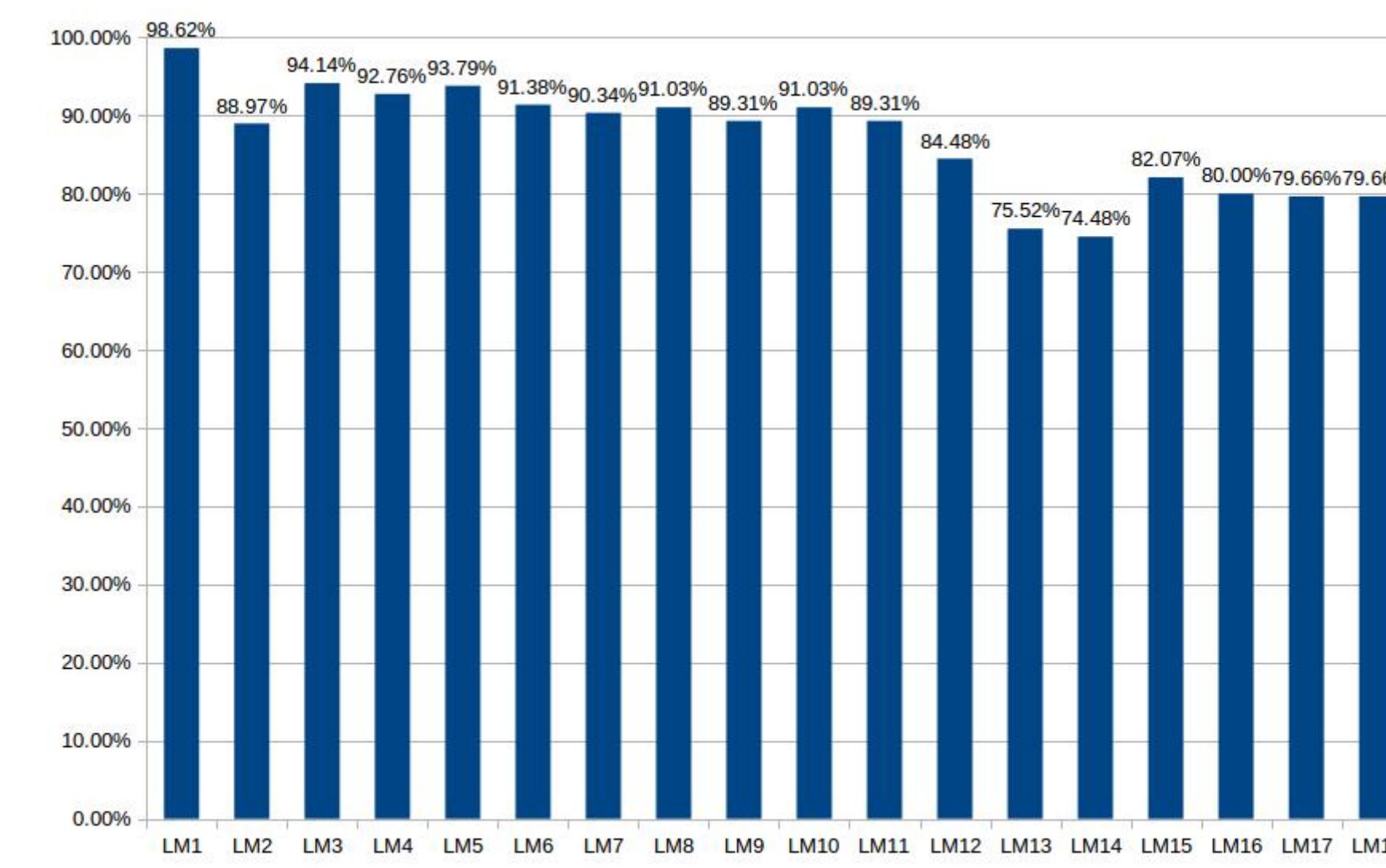
SIFT and landmarks



1. A **patch** P_s is initialized at each manual landmark of model image (size of 9×9),
2. Calculating the SIFT[4] descriptor for P_s ,
3. At the same position in the scene image, a patch P_t is created (size of 36×36),
4. For each pixel in P_t , a patch P'_t is extracted with the same size than P_s ,
5. Calculating the SIFT descriptor for all P'_t ,
6. Computing the distance between the descriptor of P_s and each P'_t ,
7. At the end, the pixel that has the **minimum distance** with P_s is kept.

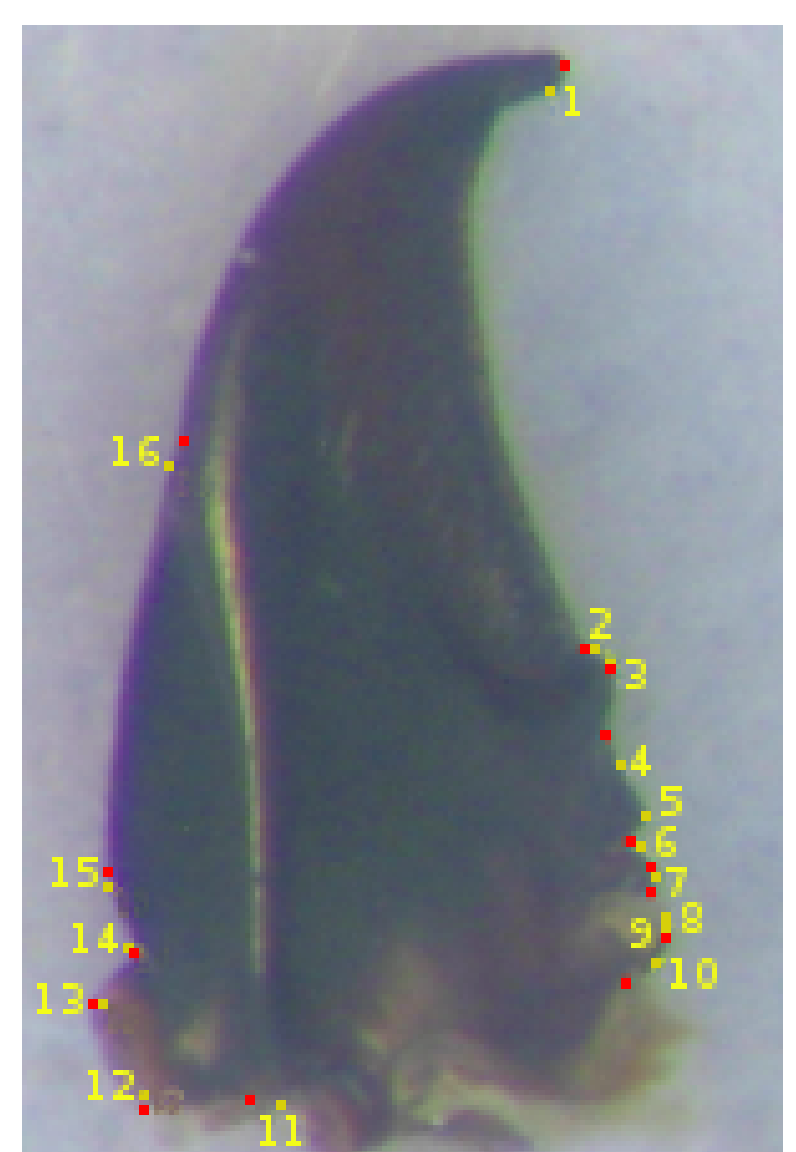
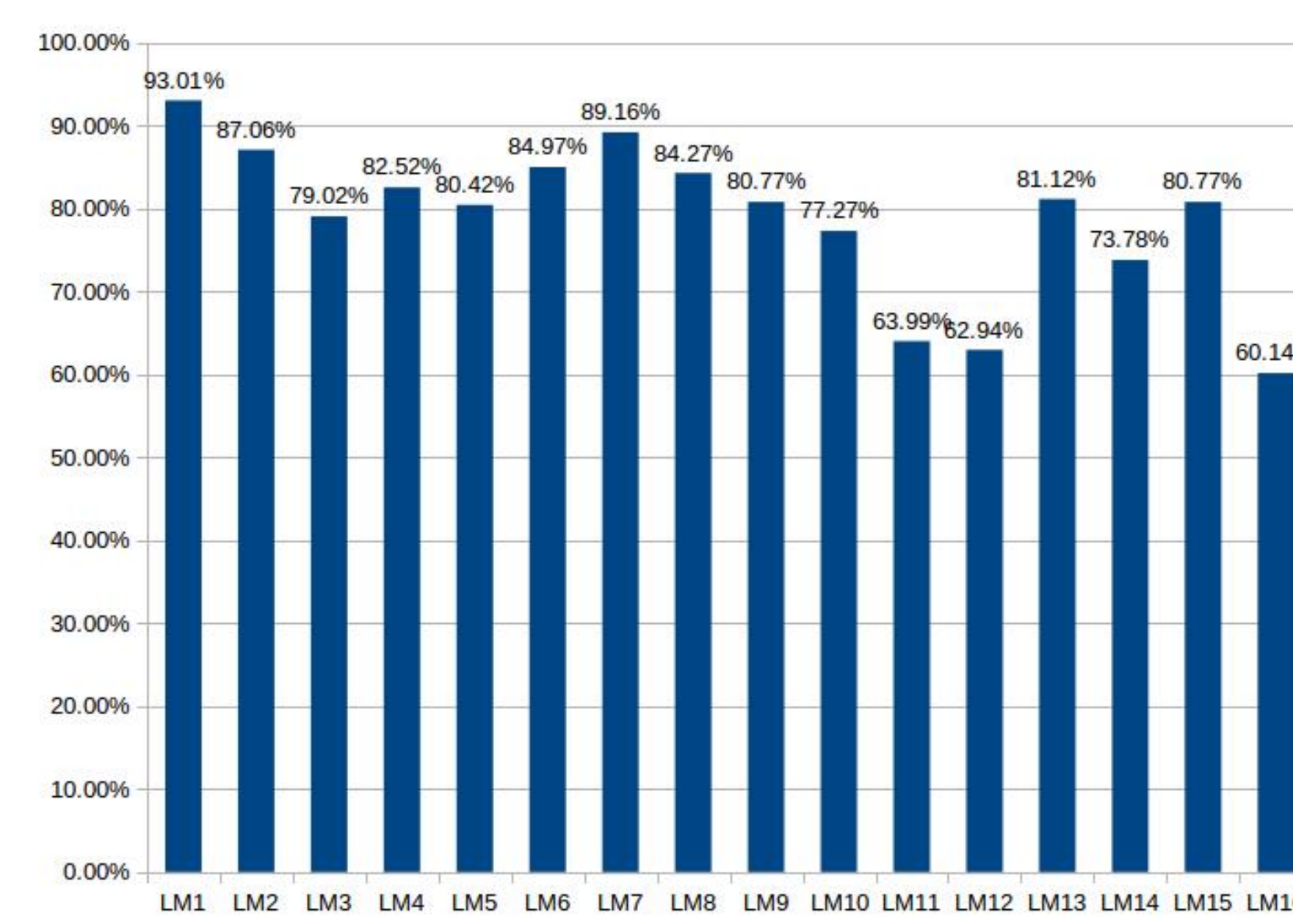
Results on right mandibles

- Highest accuracy: 1st landmark with 98.62%
- Lowest accuracy: 13th, 14th landmark with app. 75%



Results on left mandibles

- Highest accuracy: 1st landmark with 93.01%
- Lowest accuracy: 11th, 12th and 16th landmark from 60% to app. 63%



Conclusion

- A solution based on SIFT descriptor for landmark estimation is presented,
- The results show that method **succeed in locating** all landmarks in request images,
- The accuracy of method is sufficient to be **proposed to biologists** as a **replacement of manual positioning**, and to characterize the shape.

References

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