X-Ray Image Segmentation of the Hip Joint

Segmentation of the hip joint space based on a radial projection originating from the center of the femoral head

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INTRODUCTION

Osteoarthritis (OA) is a growing health problem. In 2021, 1.589.600 people were estimate to have OA in the Netherlands [1]

X-ray images are often used in the diagnosing process.

- The minimum distance between the bones in the hip joint, the minimum joint space width (JSW), is most strongly associated with OA in X-ray images [2].
- o For knee OA multiple=JSW has been proven to be even stronger associated [3].
- This might also be the case for hip OA.
- o Accurate segmentation of joint space is required to calculate the joint space profile.

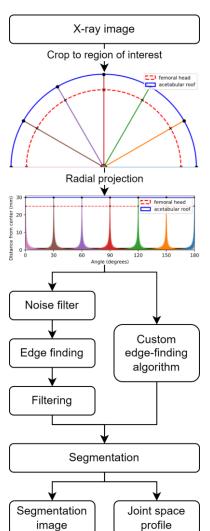
A radial projection of an image maps circular lines to straight lines.

The aim of this research was to investigate how the joint space profile can be segmented most accurately from a radial projection originating from the center of the femoral head by comparing multiple noise filtering and edge-finding algorithms.

METHODOLOGY

- Comparison of multiple denoising filters
- Comparison of multiple edge-finding algorithms
- o Development of custom edge-finding algorithm

See the figure 1 for the full pipeline.



EXPERIMENTAL SETUP

Output of algorithms are compared to landmark points created by BoneFinder [4].

Custom edge-finding algorithm:

- o Finds the strongest gradients. Those correspond to the edges of the femoral head and acetabular roof.
- Uses moving average to smoothen out outliers.

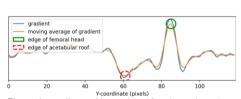


Figure 2, gradient of intensities & selected peaks

RESULTS & DISCUSSION

Best performing algorithms:

- o Custom edge-finding algorithm
- Difference of Gaussians followed by minimum cross entropy thresholding

Points of improvement:

- Roughness of edges
- o Detection start of femoral head

Can be used:

- o To measure (minimum) JSW automatically and consistently.
- In research if full joint space is a better estimator for hip OA than minimum JSW.

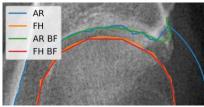


Figure 3, segmentation using custom edgefinding algorithm



Figure 4, segmentation using difference of Gaussians and minimum cross entropy thresholding



Figure 5, edges missing in segmentation

CONCLUSION

Difference of Gaussians followed by minimum cross entropy thresholding:

- o Gives reasonably good results.
- o Small blobs are removed, which are sometimes also part of the actual edge. This results in an incomplete segmentation.
- Edges are too rough

Custom edge-finding algorithm:

- Most reliable and consistent results since it always outputs an edge for the femoral head and acetabular roof.
- o Most accurate for weight-bearing part of femoral head.
- Larger errors at lateral side of femoral head due to irregularities in density and shape.
- o Start of acetabulum is often incorrect due to not having the a strong edge.

Other noise removing and edge-preserving filters do not work well with X-ray images:

- Keep too much noise.
- Smoothen out the actual edges.

[1] Vanhommerig, J., Poos, M., Gommer, A., Hendriks, C., & Giesbers, H. (2022, November). Artrose | Leeftijd en geslacht | Volksgezondheid en Zorg, Retrieved 2024-04-22, from https://www.vzinfo.nl/ artrose/leeftijd-en-geslach

[2] Croft, P., Cooper, C., Wickham, C., & Coggon, D. (1990, September). Defining osteoarthritis of the hip for epidemiologic studies. American Journal of Epidemiology, 132(3), 514-522. doi: 10.1093/oxfordjournals.aje.a115687

[3] Cheung, J. C.-W., Tam, A. Y.-C., Chan, L.-C., Chan, P.-K., & Wen, C. (2021, October). Superiority of MultipleJoint Space Width over Minimum-Joint Space Width Approach in the Machine Learning for Radiographic Severity and Knee Osteoarthritis Progression. Biology, 10(11)

[4] Lindner, C., Thiagarajah, S., Wilkinson, J. M., Wallis, G. A., & Cootes, T. F. (2013). Fully automatic segmentation of the pr random forest regression voting. IEEE Transactions on Medical Imaging, 32(8), 1462-1472. Retrieved 2024-06-10, from https://research.manchester.ac.uk/en/publications/fully-automatic-segmentation-of-the-proximal-femur-using-random-f (Publisher: IEEE) doi: 10.1109/TMI.2013.2258030

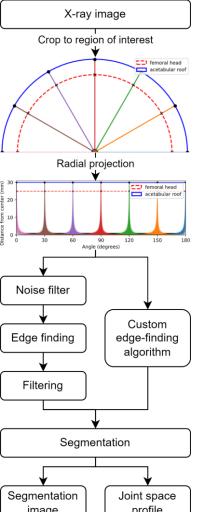


Figure 1, segmentation pipeline