

Project Part 1: Probabilistic modeling of the femur anatomy

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Abstract

The femur bone is a crucial part of the human anatomy used in forensic science due to its size and strength. It can provide valuable insights into the gender and geographical origin of the person whose remains are being analyzed. This is because femurs of people with similar geographical origin or of the same gender tend to be similar. However it is likely that when examining human remains we only get a partial or deformed femur, this is where we would need femur reconstruction to model what the full femur would have looked like and analyze it. In this project we came up with a model which we would use to reconstruct given partial femurs. We also analyzed and validated our model using statistical approaches.

Keywords: femur, forensics, gender determination, reconstruction, modeling.

1 Introduction

The main goal of this project is to develop a probabilistic shape model for femur bones which is used to reconstruct partial femurs. Femur bone reconstruction is an important task in the field of forensics as it can aid in the human identification just by analyzing the femur bones from the human remains. In this paper, we present an approach to develop a probabilistic shape model for femur bones using a combination of rigid alignment, Gaussian Process modeling, Iterative Closest Points (ICP), and Principal Component Analysis (PCA).

In our approach we used a dataset of femur meshes and femur landmarks, comprising 47 samples each. We first applied rigid alignment, followed by building a Gaussian Process model (GP-model) to capture the shape variation of the femur bones. We then used an iterative alignment refinement technique called ICP to fit the GP-model by aligning the femur meshes with the model.

After implementing ICP, we built a PCA model to reduce the dimensionality of the data and speed up the reconstruction process.

2 Methods

2.1 Rigid Alignment

2.2 GP model

2.3 ICP

2.4 PCA

In this section you can describe the methods you used to solve the problem.

3 Experiments and results

3.1 Data and experimental setup

The data on which we are experimenting consists of 46 femurs. Included in this data are landmarks L_0, \dots, L_5 for every femur. In particular L_3, L_4 , as seen in figure 1, can be used to estimate the width of the femur, whereas L_2, L_5 , as seen in figure 2, can be used to estimate the length.

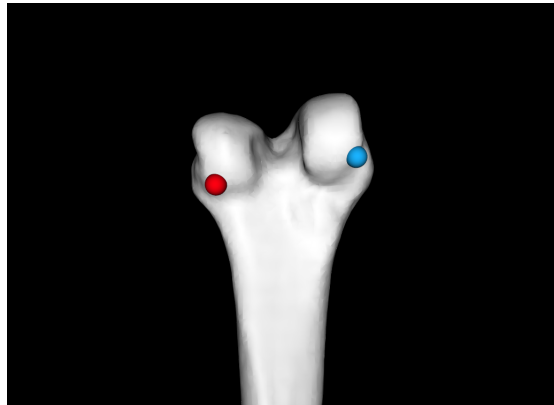


Figure 1: Landmark L_3 in blue, and landmark L_4 in red. Used to estimate width of the femur bone.

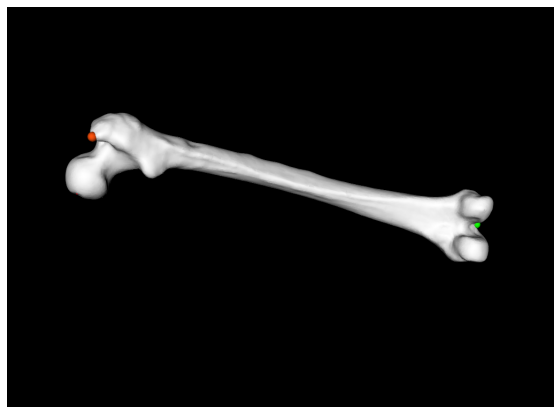


Figure 2: Landmark L_3 in blue, and landmark L_4 in red. Used to estimate length of the femur bone.

To estimate the length of a femur we calculate the distance between the L_2 and L_5 landmark, similarly we estimate the width by calculating the distance between the L_3 and L_4 landmark. We display the measurements made from the 46 femurs in figure 3 and summarize them in table 1 by calculating the mean and variance.

	Mean	Variance
Length	420.80	841.94
Width	60.61	18.13

Table 1: Mean and variance of length and width of the femur data.

We notice a high variability of the data. This gives reason to believe that the 46 femur samples are not from subjects that all share similar femurs. More precisely, the data probably originates

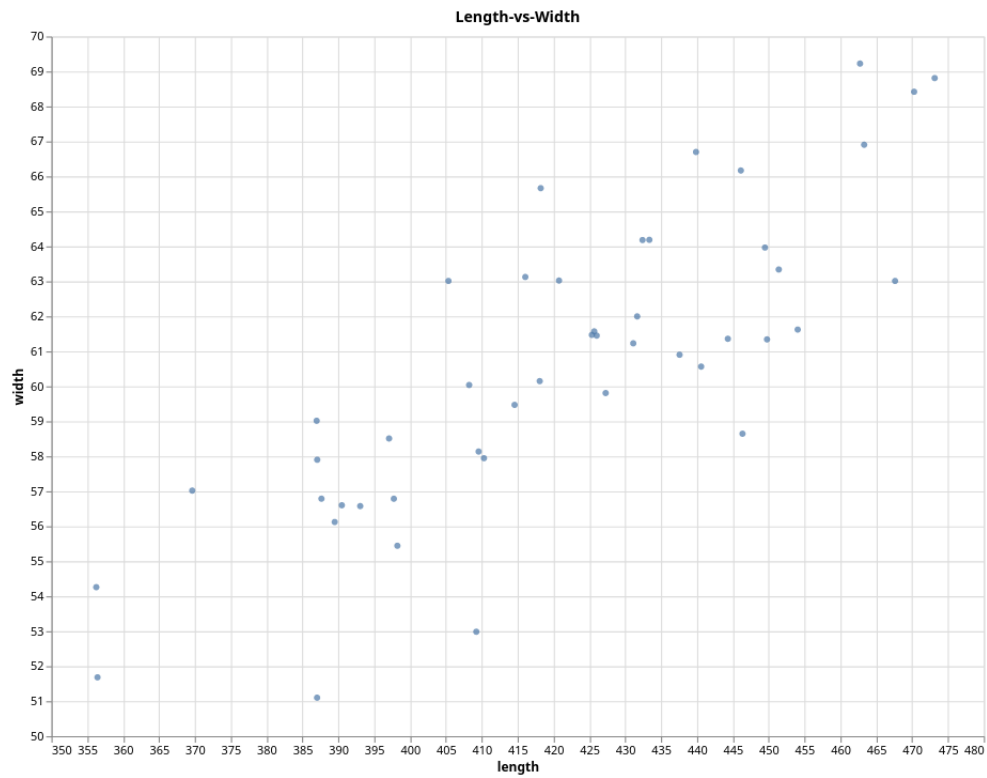


Figure 3: Scatterplot of calculated length and width of the femur data

from males and females with a large variability in height, and therefore, a large variability in femur length and width. Visually, figure 3 gives reason to believe that the length and width are correlated. There is a trend that shows that an increase in length comes with an increase in width.

3.2 Experimental results

4 Conclusion

Add your conclusion here. What is the main result? What did you achieve, what needs to be done.