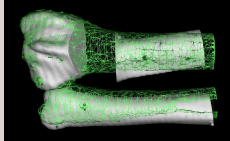


I. Introduction

Motivation

- A surgeon must manually align models of a damaged radius (white) and the mirrored radius of the opposite limb (green) in the Distal Radius Osteotomy (DRO) planner of Croitoru et. al. [2]

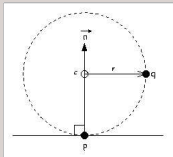


- Aligning models in 3D is time consuming and frustrating for the surgeon
- Assuming both arms are roughly symmetric, a shape-based registration algorithm such as ICP[1] might be used to speed up or even automate the alignment procedure
- We examine using ICP to align the damaged distal fragment (after corrective cut) with the mirrored radius of the opposite limb

Hypotheses

- On average, ICP produces alignments with small angular and lengthening deviations from those planned by a surgeon
- Using registration points of *high* curvature produces smaller deviations from surgeon-planned alignments than using all available points or points of *low* curvature

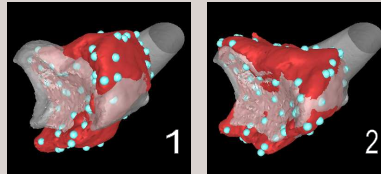
Minimal Local Curvature at a Point



The inverse of the radius (r , in mm) of the largest circle in the set of circles containing a point (p) and each neighbour (q) of p . Circle centers are located along the point normal (n) of p . Circles lie in the plane defined by p , q , and c [3].

Registration Point Selection Factors

- Number of points (50 vs. 100)
- Random selection vs. subsampling
- Point set
 - All points
 - High curvature (min. local curv. $\geq 1/5$)
 - Low curvature (min. local curv. $< 1/5$)
 - Equal number of high and low curvature



Experimental Procedure

- Experimental Trials:
 - Registration points selected from affected distal fragment
 - ICP applied to align affected distal fragment with reflected undamaged radius
- For each test scene: one trial for subsampling conditions, 1000 trials for random selection conditions
- Factorial Design: all combinations of point selection factors above examined

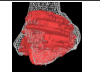


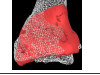



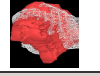

Metrics Observed After Alignment:

- Angle between long axes in anterior-posterior view (AP)
- Angle between long axes in mediolateral view (ML)
- Angle between palmar-dorsal axes in axial view (AX)
- Distance between furthest affected and unaffected points along the long axis of the unaffected radius (Length)

Qualitative Evaluation

- The surgeon that performed the original procedures visually inspected the ten results for the *subsampling 50 of all points* condition
- The surgeon classified six results as usable as-is, three as requiring minor alteration, and one as requiring 'major' alteration

Table 2. Sample ICP Distal Radius Alignment Results¹

Informal Classification	Anterior-Posterior View	Mediolateral View	Axial View
Usable As-Is			
Minor Alteration			
Major Alteration			

IV. Discussion

Points of High Curvature

- Likely include spurs and other artifacts on the affected distal fragment

Small Lengthening Deviations

- Possibly due to the initial translation of the affected distal fragment during test-scene creation

Experimental Design Limitations

- Used a very simple curvature metric; others may be worth studying
- Low power (only ten test scenes)

Using Multiple Initial Alignments

- Better to offer a surgeon a small set of initial alignments rather than a single one in practice

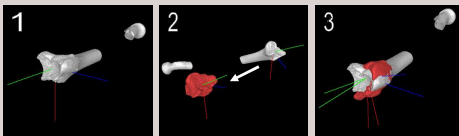
II. Methods and Materials

Ten Ground-Truth Test Scenes

From DRO plans previously executed by a single surgeon

Ground-Truth Procedure

- Axes placed manually on reflected unaffected radius (long (green), lateral (blue), palmar-dorsal (red) axes)
- Axes mapped onto damaged distal fragment using inverse of corrective transformation in original surgical plan
- Distal fragment manually translated roughly into place



III. Results

Table 1. Deviations of ICP Results from Original DRO Plan (10 Test Scenes, Subsampling Condition⁵)

Number Points	AP ¹ (degrees)	ML ² (degrees)	AX ³ (degrees)	Length ⁴ (mm)
100 points				
mean, std	3.2, 2.6°	6.7, 5.9°	4.5, 5.0°	1.08, 1.06 mm
range	0.2 – 7.8°	1.7 – 19.2°	0.5 – 13.1°	0.01 – 3.24 mm
50 points				
mean, std	6.5, 4.8°	10.8, 9.1°	4.3, 4.8°	1.03, 0.78 mm
range	0.1 – 14.4°	1.5 – 27.1°	0.1 – 14.5°	0.18 – 2.75 mm

¹AP: angular difference of long axes in anterior-posterior view
²ML: angular difference of long axes in mediolateral view
³AX: angular difference of palmar-dorsal axes in axial view
⁴Length: length difference along long axis of model bone
⁵Subsampling Condition: ICP using registration points subsampled from all affected distal fragment surface model points

ANOVA Results

- Two-way ANOVA performed for each metric (sampling, point set factors combined)
- Means of random results used to create a balanced design
- Main effect for number of points detected for the AP metric ($p < 0.05$). The mean AP value for the 100 points group (3.9°) was 1.1° smaller than the mean for 50 points (5.0°)

V. Conclusion

Hypotheses

- Mean angular and lengthening deviations of ICP alignments from surgeon-planned alignments were small
- Using registration points of high curvature *did not* reduce deviation from the surgeon-planned alignments

Visual inspection of results indicates that ICP is a useful addition to computer-assisted DRO planning

References

- Bestl and McKay. IEEE TPAMI 1992; 14(2): 239-256.
- Croitoru, Ellis, Small and Pichora. LNCS 2000; 1935:1126-1135
- Schneider and Kobbelt. Computer-Aided Geometric Design 2001; 18:359-379.