

# OpenArm 2.0: Automated Segmentation of 3D Tissue Structures for Multi-Subject Study of Muscle Deformation Dynamics

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## Contributions

- Novel **neural-network-based pipeline for segmentation of 3D muscle and bone structures** from localized 2D ultrasound data of the human arm
- The **OpenArm 2.0 data set**, the first factorial set of **multi-subject, multi-angle, multi-force scans of the arm with full volumetric annotation** of the biceps and humerus

## Target Application Domains

- **Musculoskeletal simulation:** measurement of individual muscle forces
- **Assistive device control:** extraction of multiple robust control signals for high-DoF prosthesis / exoskeleton control
- **Diagnosis / rehabilitation:** improved measurement and understanding of musculoskeletal deficiency
- **Graphics / animation:** enhanced rendering of muscle shape changes during movement

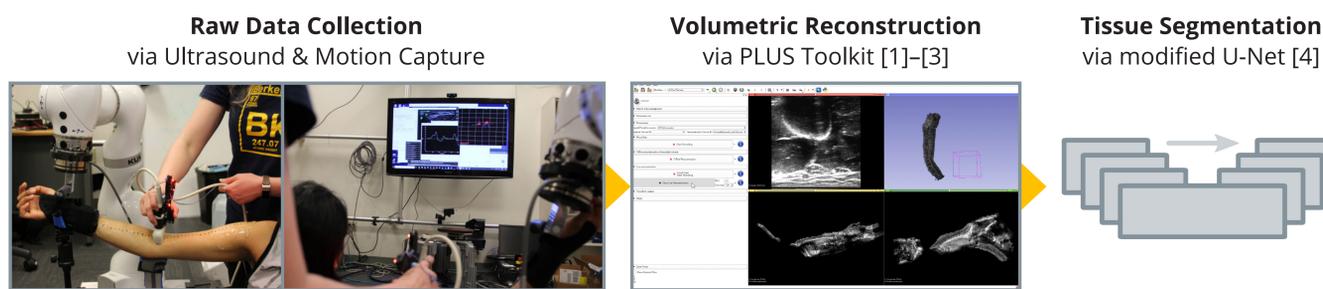
## DOWNLOAD

Download all code and data at  
simtk.org/projects/openarm



## Data Set Collection & Specifications

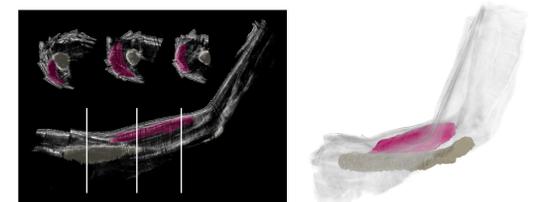
### Final Data Processing Pipeline



Full 3D intensity maps of the anterior surface of the arm were assembled from 2D ultrasound data (localized via motion capture) using an adapted version of the first OpenArm experimental protocol [5].

Several improvements were made, including **real-time force tracking and visual feedback** to enable scan collection under arbitrary, repeatable force conditions.

### Finalized Tissue Volumes



Data were collected from **10 subjects** (+1 partial) at

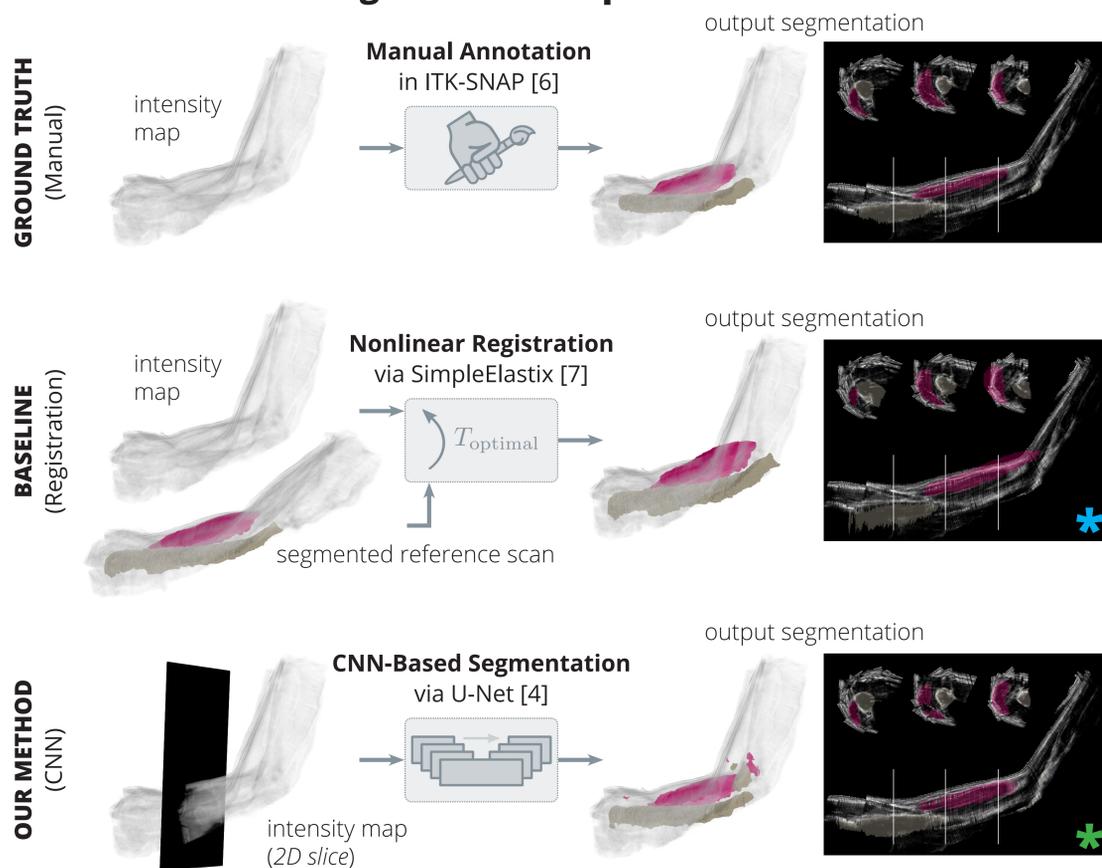
- 4 elbow angles (0°, 30°, 60°, 90°)
- 5 force conditions ("fully supported", 0/10/30/50% MVC)

**20 scans / subject**

**automated annotation** performed for all scans  
**ground truth annotation** performed for 1 full + 2 partial subject data sets

## Automated Tissue Segmentation

### Segmentation Pipelines



### Segmentation Accuracy by Strategy

Architecture / Strategy	same angle, same force, same subject	Accuracy (IoU, Pixel Accuracy)			
		new angle, same force, same subject	same angle, new force, same subject	same angle, same force, new subject	same angle, same force, new subject
BASELINE (Registration)	RR	0.431, 0.980	0.320, 0.970	0.537, 0.985	0.244, 0.953
	RANR	0.722, 0.991	0.545, 0.980	0.450, 0.980	0.386, 0.960
OUR METHOD (CNN)	U-NET	0.875, 0.996	0.593, 0.982	0.604, 0.988	0.422, 0.961
	U-NET+RA	0.929, 0.998	0.560, 0.984	0.464, 0.986	0.393, 0.971
	U-NET+EA	0.950, 0.999	0.677, 0.988	0.573, 0.989	0.533, 0.978
	U-NET+RA+EA	0.936, 0.998	0.577, 0.984	0.544, 0.988	0.499, 0.978
	Multi-Angle U-NET	0.886, 0.997	0.691, 0.989	0.614, 0.989	0.470, 0.972
	Multi-Angle U-NET+EA	0.906, 0.997	0.717, 0.989	0.651, 0.990	0.523, 0.975
	Multi-Force U-NET	0.885, 0.997	0.617, 0.985	0.770, 0.994	0.452, 0.972
	Multi-Force U-NET+EA	0.902, 0.997	0.682, 0.988	0.793, 0.994	0.519, 0.977
	Multi-Subject U-NET	0.884, 0.997	0.657, 0.987	0.536, 0.988	0.885, 0.995
	Multi-Subject U-NET+EA	0.908, 0.998	0.687, 0.989	0.565, 0.989	0.909, 0.996

Training Accuracy: "memorization baseline" (grey bar)  
Validation Accuracy: BEST (purple) to WORST (orange)

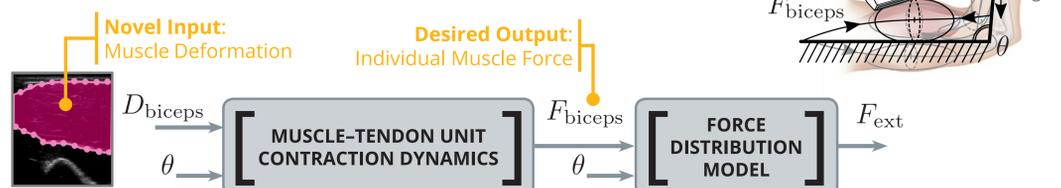
Our CNN-based segmentation method results in **more accurate tissue segmentation than baseline registration-based methods**, especially when networks are trained on **elastically augmented data** [8] from **multiple subjects**.

Registration-based methods result in significant errors **along the entire length of the muscle**, while our CNN-based methods reliably segment the muscle belly, with errors **primarily at muscle ends**. Our method thus results in **faster manual cleanup time** when it is necessary for applications requiring higher accuracy.

Segmenting new subjects remains more difficult than segmenting new angle and force conditions; thus, we are **actively working to improve accuracy** by training **subject-specific networks on more comprehensive data sets**.

## Current Work: Muscle Force Modeling

**CORE HYPOTHESIS:** Individual muscle force can be inferred from muscle deformation, which can be detected via ultrasound. This relationship can be measured and quantified because changes in muscle shape reflect changes in tendon length, and therefore tendon stiffness, the mechanism by which force is imparted to the skeleton.



We are building a **principled suite of models** that make varying trade-offs between **collected data** and (possibly unreliable) **literature values** in a quantifiable manner, ranging from "black box" to "white box".

## Acknowledgments / Sponsors / References

The authors acknowledge the aid and advice of **Dr. Gregorij Kurillo**, **Akira Kato**, **Jaeyun Stella Seo**, **David Wang**, **Sachiko Matsumoto**, and **Nandita Iyer** in both hardware development and data collection; of **Jeffrey Zhang** and **Kireet Agrawal** in development of initial U-Net and image processing code; and of **Ian McDonald** in development of image registration code.

This work was supported by the **NSF National Robotics Initiative** (award no. 81774), **Siemens Healthcare** (85993), the **NVIDIA Corporation GPU Grant Program**, **eZono AG**, and the **NSF Graduate Research Fellowship Program**.

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