

OpenKnee Data Navigation - Registration

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Download Data

- Joint mechanics
- Registration marker STLs
- Setup knee directory
 - geometry-oks0XX: contains bone STLs built from segmentation (in repository)
 - joint_mechanics-oks003: contains all joint mechanics data (downloads section of SimTk)

SimTK

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ok Open Knee(s): Virtual Biomechanical Representations of the Knee Joint

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Open Knee(s) - Generation 2 - Specimen 1

g2-s1-v0.2.0.20150825

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1.3.12.2.1107.5.2.19.45406.2014100711193292568244326.0.0.0.nii	AUG 24, 2015
21 MB Any Data/Images/Video	
Soft tissue imaging - axial plane (MRI in NIFTI format)	
1.3.12.2.1107.5.2.19.45406.2014100710433217692143626.0.0.0.nii	APR 16, 2015
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Cartilage imaging (MRI in NIFTI format)	
1.3.12.2.1107.5.2.19.45406.2014100711262396541244530.0.0.0.nii	APR 16, 2015
21 MB Any Data/Images/Video	
Soft tissue imaging - sagittal plane (MRI in NIFTI format)	
1.3.12.2.1107.5.2.19.45406.2014100711323578731244734.0.0.0.nii	APR 16, 2015
21 MB Any Data/Images/Video	
Soft tissue imaging - coronal plane (MRI in NIFTI format)	
1.3.12.2.1107.5.2.19.45406.2014100710220888421542662.0.0.0.nii	APR 15, 2015
92 MB Any Data/Images/Video	
General purpose imaging (MRI in NIFTI format)	
joint_mechanics-oks001.zip	AUG 24, 2015
226 MB Any Data/Images/Video	
Joint experimentation files for characterization of tibiofemoral and patellofemoral joint mechanics (text and binary file formats)	

joint_mechanics-oks001.zip

226 MB Any Data/Images/Video

Joint experimentation files for characterization of tibiofemoral and patellofemoral joint mechanics (text and binary file formats)

AUG 24, 2015

Setup for registration

- Make sure there is a file with digitizing order within the main Configuration folder of the tibiofemoral joint testing directory (ReadMe_Notes.txt)
- Create registration XML
 - Root directory of knee data
 - Left or Right?
 - Bone STLs
 - Sphere STLs

MRI Spheres

Tibia:

Sphere 1 = Lateral
Sphere 2 = Medial
Sphere 3 = Posterior

Femur:

Sphere 1 = Lateral
Sphere 2 = Medial
Sphere 3 = Posterior

```
<Registration_Input>
  <Knee_directory>/home/morille2/Documents/OpenKnee/oks001</Knee_directory>
  <Knee_side>Right</Knee_side>
  <Bones>
    <Femur>
      <file>oks001_MRG_FMB_CPL_RAW_02.stl</file>
    </Femur>
    <Tibia>
      <file>oks001_MRC_TBB_CPL_RAW_02.stl</file>
    </Tibia>
    <Fibula>
      <file></file>
    </Fibula>
    <Patella>
      <file>oks001_MRG_PTB_CPL_RAW_02.stl</file>
    </Patella>
  </Bones>
  <Registration>
    <FMR>
      <Posterior>oks001_MRG_FMR-P_RAW_02.stl</Posterior>
      <Lateral>oks001_MRG_FMR-L_RAW_02.stl</Lateral>
      <Medial>oks001_MRG_FMR-M_RAW_02.stl</Medial>
    </FMR>
    <TBR>
      <Posterior>oks001_MRG_TBR-P_RAW_02.stl</Posterior>
      <Lateral>oks001_MRG_TBR-L_RAW_02.stl</Lateral>
      <Medial>oks001_MRG_TBR-M_RAW_02.stl</Medial>
    </TBR>
    <PTR>
      <Superior>oks001_MRG_PTR-S_RAW_01.stl</Superior>
      <Lateral>oks001_MRG_PTR-L_RAW_01.stl</Lateral>
      <Medial>oks001_MRG_PTR-M_RAW_01.stl</Medial>
    </PTR>
  </Registration>
</Registration_Input>
```

Coordinate Systems

$$P_B = (T_{A_B})^{-1} \cdot P_A$$

Transformation of points from A CS to B CS – T_{A_B} .

Transform B coordinates in the A coordinates.

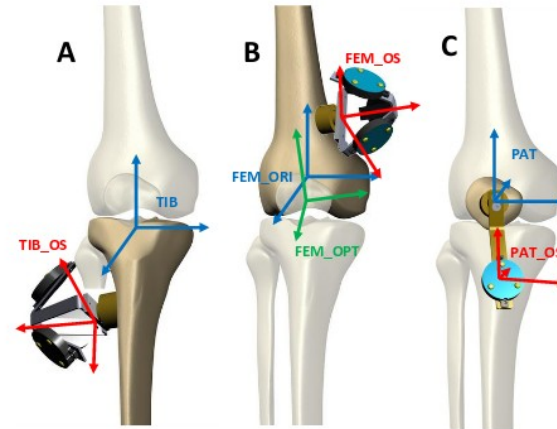
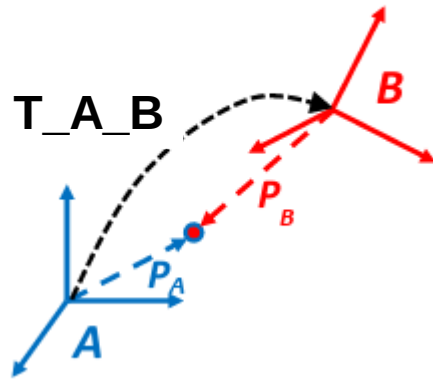


Figure 3: Coordinate Systems for A) tibia, B) femur and C) patella

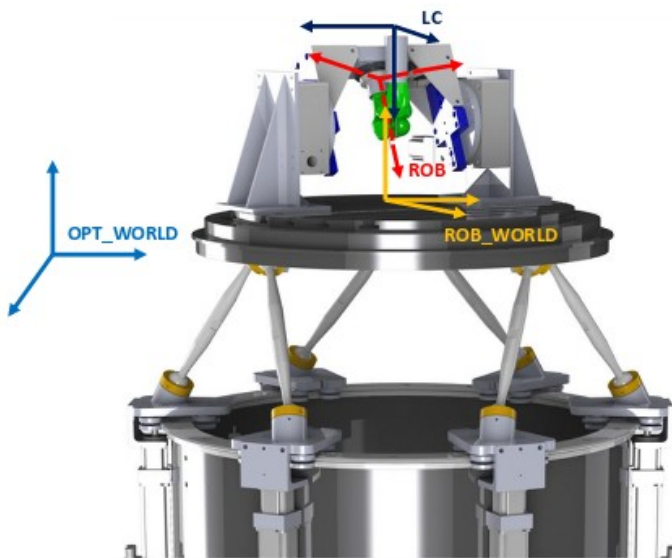


Figure 1: Coordinate Systems for Tibiofemoral testing

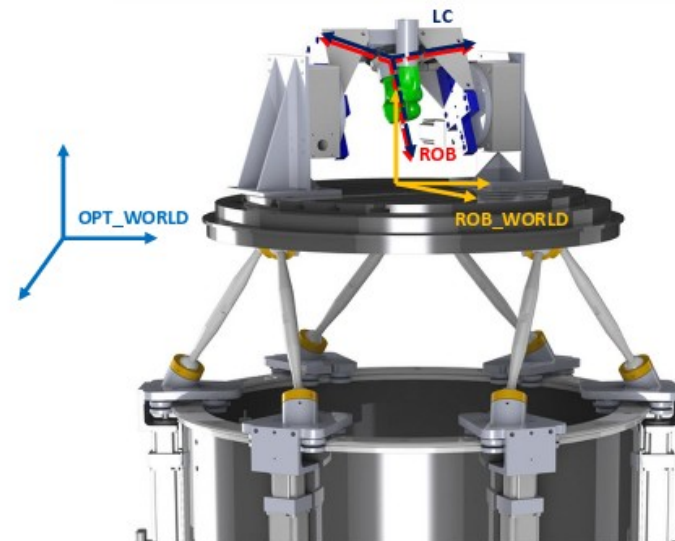


Figure 2: Coordinate Systems for Patellofemoral Testing (Note: During patellofemoral testing, the tibia is mounted to the robot and the femur is mounted to the frame. Also, there are two physical load cells attached to the stage that are not displayed in the image)

Registration Script

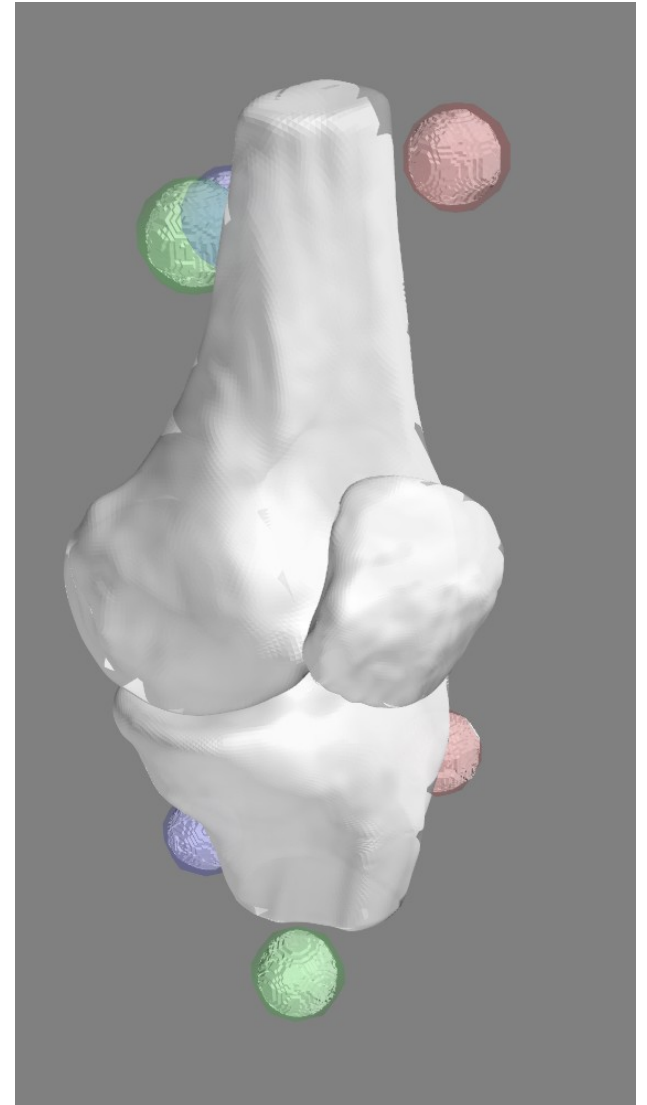
- Change directory to desired OpenKnee specimen
- Run `convert_to_image_CS.py <oks0XX_registration_0x.xml> <1>`
 - Inputs:
 - Registration XML
 - Boolean: 0 – no visualization, 1 – visualization
 - Outputs:
 - Transformation matrices saved within .npz file for each trial with processed TDMS data.

What is the script doing?

- **Performing patella, tibia, and femur registration using digitized points and segmented spheres captured in MRI.**

Tibiofemoral Registration

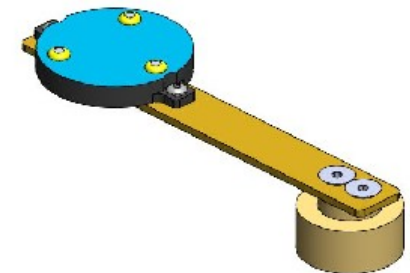
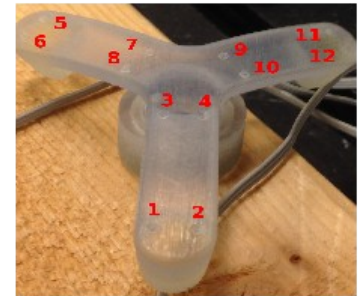
- Completed for each individual TF joint trial
- Transformations directly from the State.cfg file
 - `T_TIBOS_TIB`
 - `T_FEMOS_FEMORI`
 - `Offset_FEMORI_TIB`
- Extract sphere centers
 - Tibia and Femur: 3 spheres per bone
 - Two coordinate systems to relate experiment to image/model coordinate systems
 - Digitized during experimentation
 - Points are collected in the WORLD CS, but position sensor data is also collected for the relevant bone. Transform these points into FEMOS or TIBOS.
 - Fit sphere to digitized points and find the center
 - STLs from MRI segmentation
 - Fit sphere to STL surface to get center (image coordinate system)
 - Singular value decomposition (SVD) to find `T_I_TIBOS` and `T_I_FEMOS`



Patella Registration

- Extract divot points that were digitized on the registration marker assembly (State.cfg file of patella registration folder)
 - Points converted to REFOPT CS
 - $T_{\text{REFOPT_WORLD}}$
- Extract CAD points (CAD_PT_DIMENSIONS.txt)
 - Divots
 - Spheres
- Extract sphere centers (STLs in MRI)
- $T_{\text{REFOPT_PATOS}}$ (Registration marker relative placement.tdms)
 - $T_{\text{WORLD_PATOS}}$ from patella sensor
 - $T_{\text{WORLD_REFOPT}}$ from reference sensor
- $T_{\text{REFOPT_CAD}}$: SVD of divots in CAD and REFOPT CS
- $T_{\text{I_CAD}}$: SVD of sphere centers from CAD and MRI
- $T_{\text{I_PATOS}}$: built from the transformation matrices above.
 - Note: This registration is only performed once for all trials!

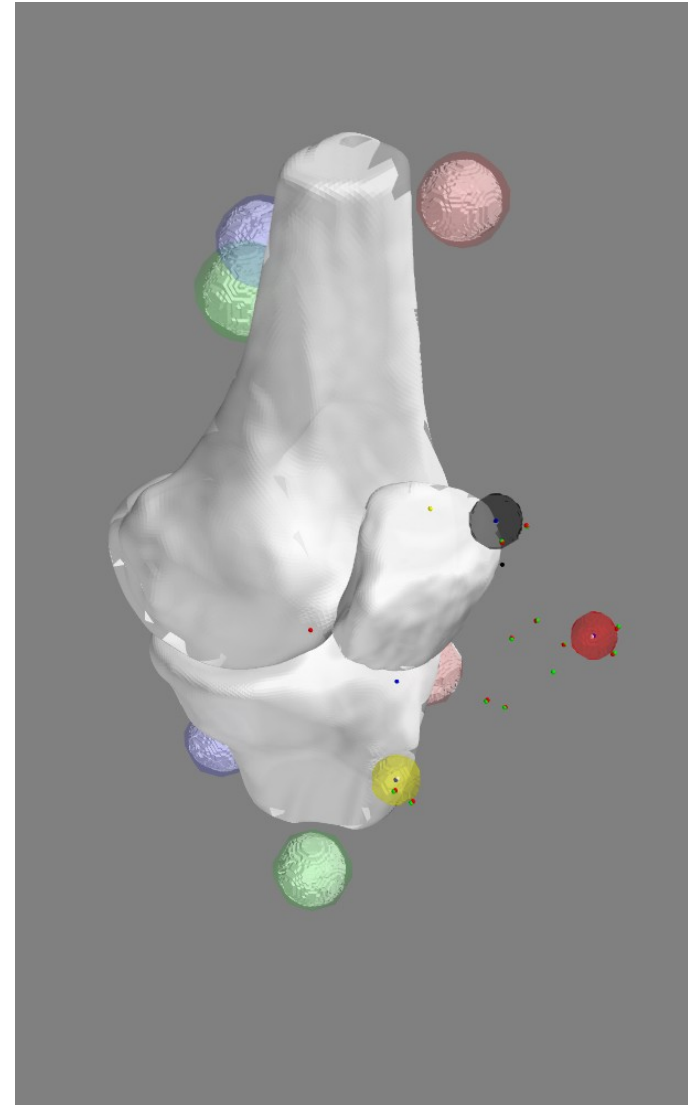
Registration marker assembly



Patella Optotrak sensor

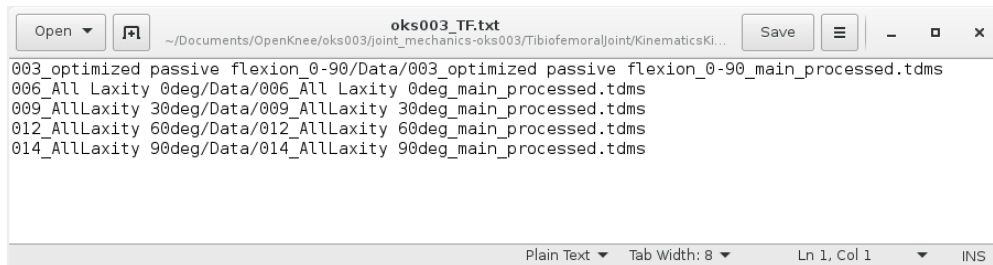
Patellofemoral Registration

- Completed for each individual PF joint trial
- Transformations directly from the State.cfg file
 - T_TIBOS_TIB
 - T_FEMOS_FEMORI
 - T_PATOS_PAT
 - Offset_FEMORI_TIB
 - Offset_FEMORI_PAT
- Extract sphere centers (same process as TF joint trials)
 - T_I_TIBOS and T_I_FEMOS



Re-sampling Data Script

- After registration script has been run.
- Run `resampling_plotting.py <oks0XX_registration_0x.xml> <tdms_list.txt>`
 - Inputs:
 - Registration XML
 - Text file of tdms files that you would like to re-sample (text file needs to be within the general test folder, i.e. TibiofemoralJoint or PatellofemoralJoint)



- Outputs:
 - CSV file of all channels within the TDMS file at the re-sampled locations
 - Video animation of the test
 - Graphs of the JCS kinematics and loads (both raw and re-sampled)

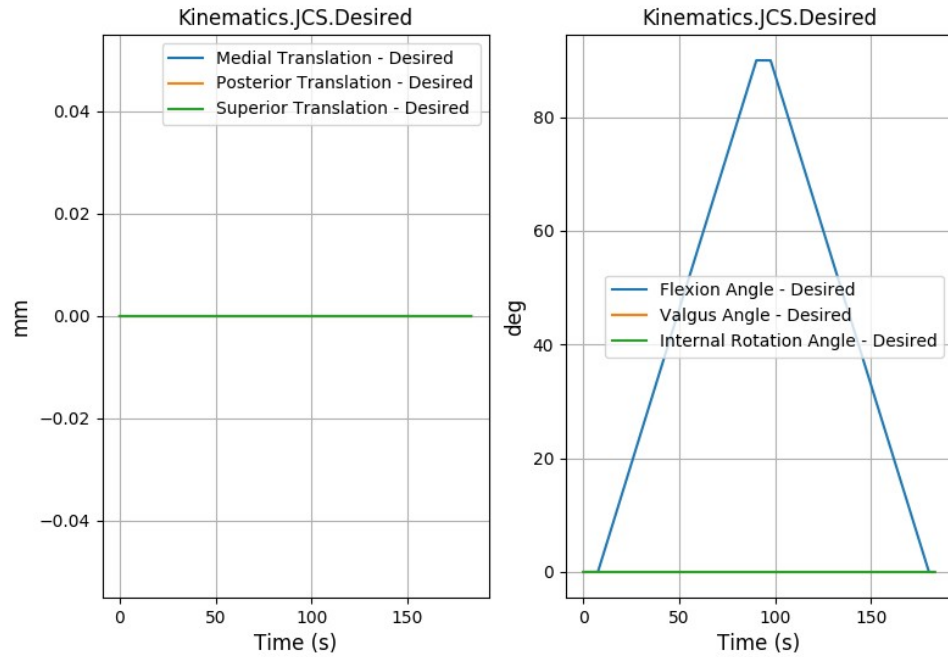
What is the script doing?

- **Sampling the data into a more usable format for comparison to model results and/or boundary condition input.**

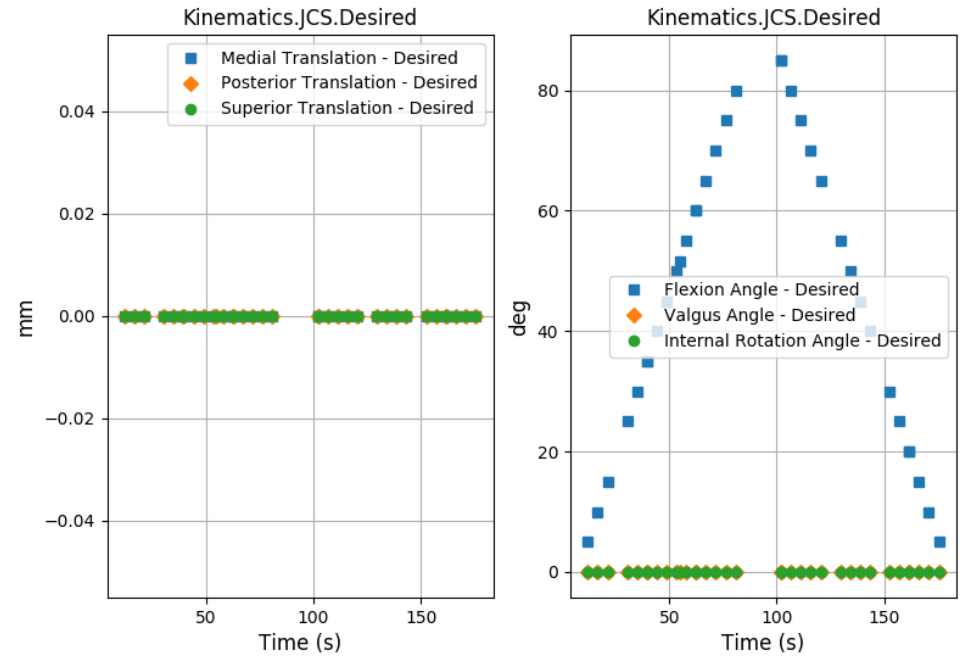
Animation – Passive flexion oks003



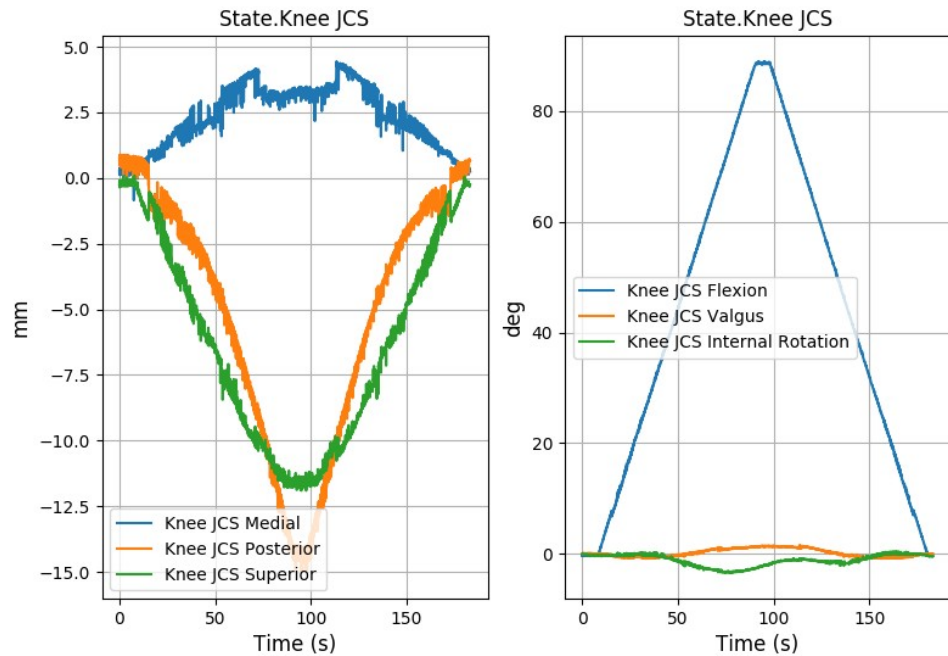
003_optimized passive flexion_0-90_main_processed



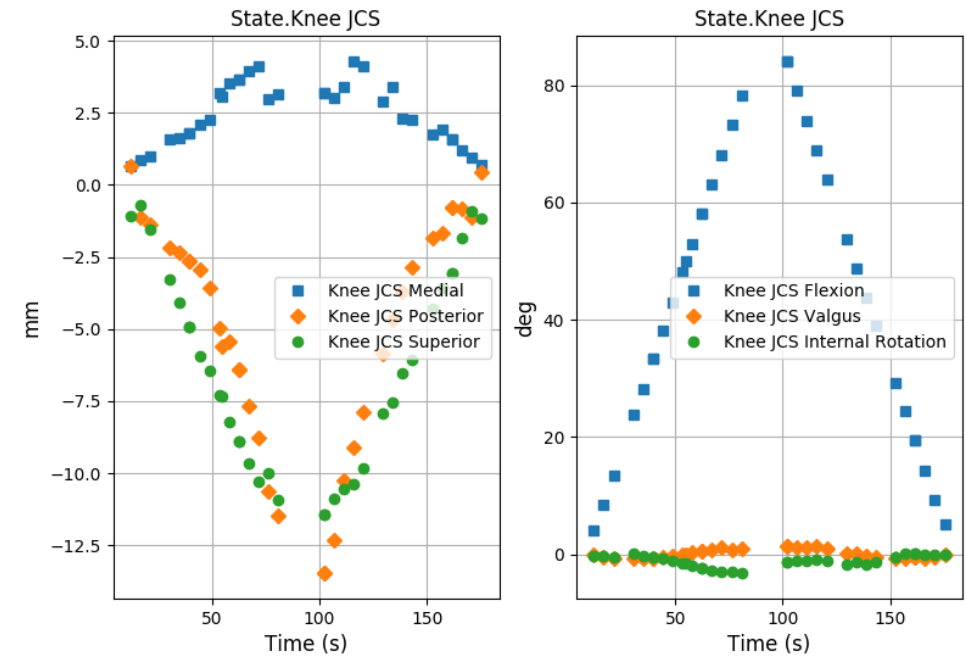
003_optimized passive flexion_0-90_main_processed_extracted



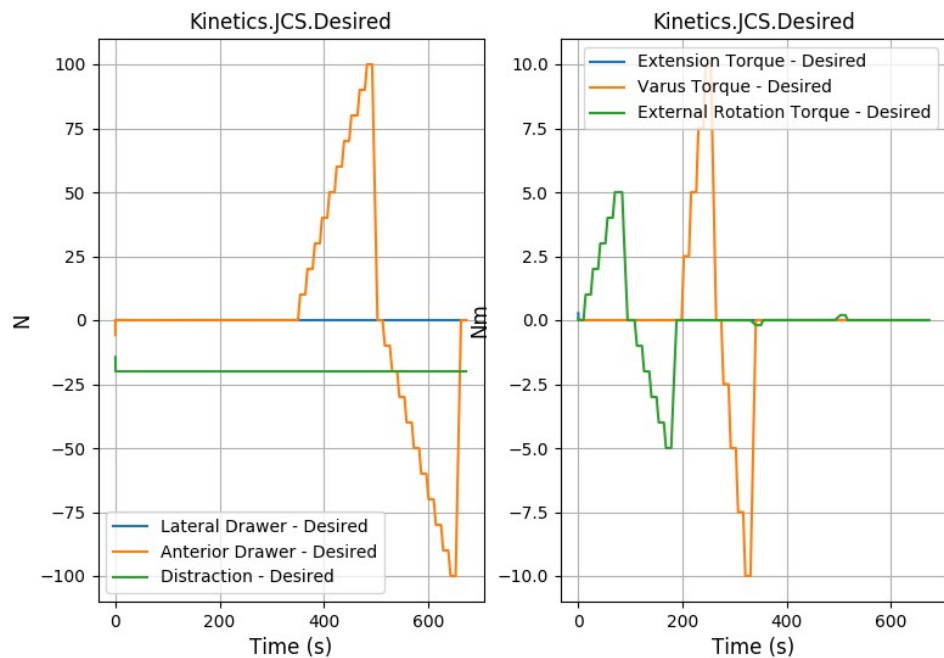
003_optimized passive flexion_0-90_main_processed



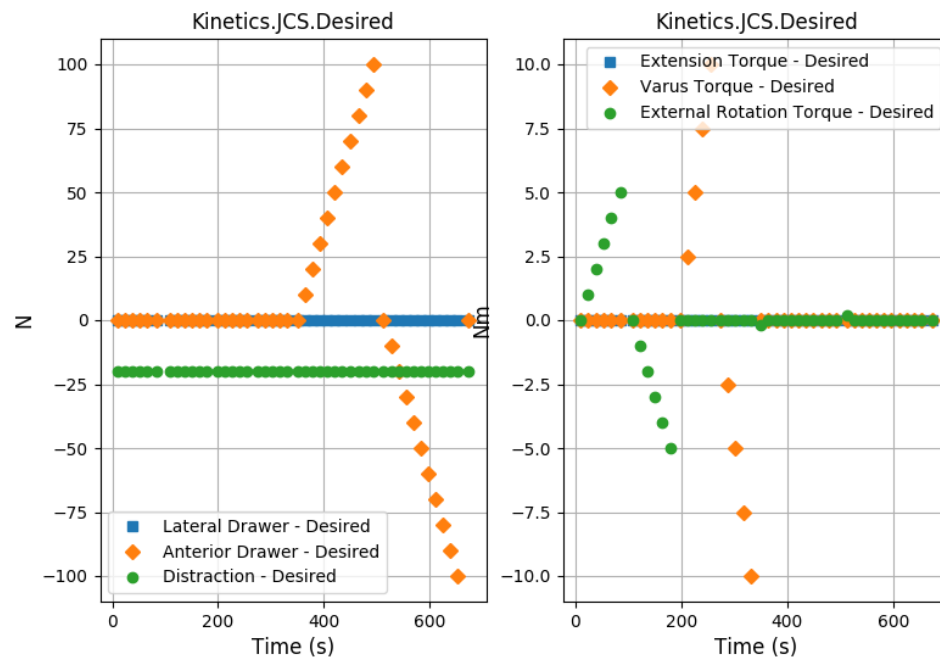
003_optimized passive flexion_0-90_main_processed_extracted



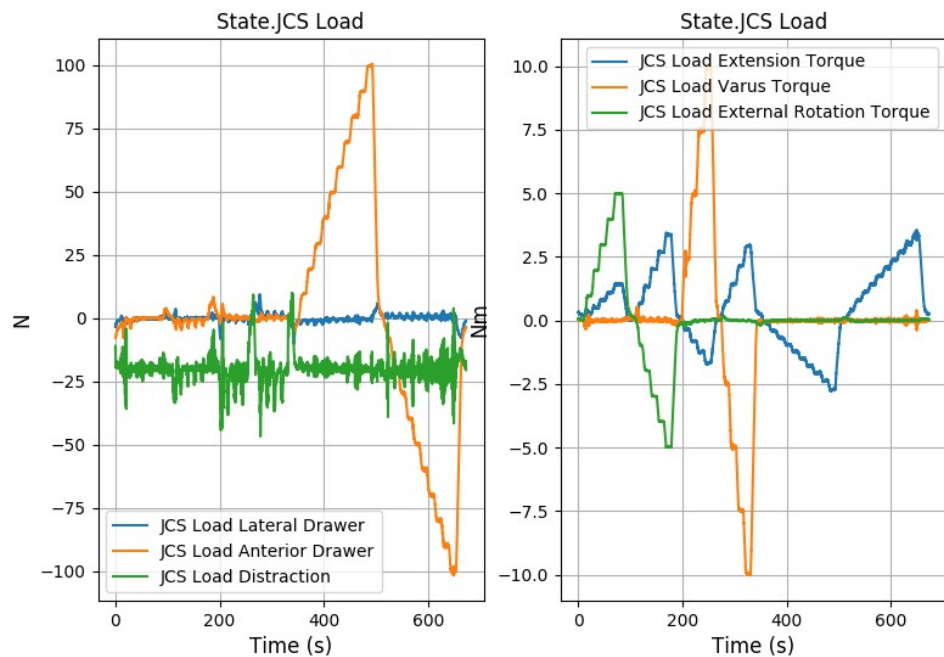
006_All Laxity 0deg_main_processed



006_All Laxity 0deg_main_processed_extracted



006_All Laxity 0deg_main_processed



006_All Laxity 0deg_main_processed_extracted

