



In biomechanical research, we often collect motion capture (marker trajectories) and force (analog data) data to calculate joint kinematics (joint angles and joint angular velocities) and joint kinetics (joint moments and joint power). Usually, this experimental data is collected with motion capturing software such as Qualysis Track Manager or Vicon Nexus, to name only a few. Regardless of the software used to capture the biomechanical data, the C3D file extension has been established as a standard for further data processing. Although C3D-files can be opened by the Motion Capturing Software, resource-saving and open source tools are available to visualize and manipulate C3D files. In this tutorial you will learn how to

1. [Visualize C3D files using Mokka](#)
2. [Load C3D files with BTK in Matlab](#)
3. [Downloads](#)

1. Visualize C3D files using Mokka

Mokka is free to use and cross-platform software to easily analyze biomechanical data. Mokka can be downloaded from the website <http://biomechanical-toolkit.github.io/mokka/index.html> and requires no installation.

After downloading and unpacking the Mokka version for your operating system, the folder should look like figure 1. Click on Mokka and a graphical user interface (GUI) that looks like figure 2 will open.

MokkaHelp	15.02.2013 16:59	Dateiordner	
phonon_backend	14.12.2012 09:40	Dateiordner	
Mokka	06.05.2013 11:02	Anwendung	9.018 KB
MokkaUpdater	02.05.2013 10:34	Anwendung	33 KB
msvc90.dll	16.04.2012 09:13	Anwendungserwei...	834 KB
msvcr90.dll	16.04.2012 09:13	Anwendungserwei...	627 KB
phonon4.dll	03.12.2012 13:52	Anwendungserwei...	334 KB
QtCore4.dll	03.12.2012 13:01	Anwendungserwei...	3.121 KB
QtGui4.dll	03.12.2012 13:31	Anwendungserwei...	10.412 KB
QtNetwork4.dll	03.12.2012 13:04	Anwendungserwei...	1.140 KB
QtOpenGL4.dll	03.12.2012 13:38	Anwendungserwei...	897 KB

Figure 1: Mokka folder after unpacking the .zip file.

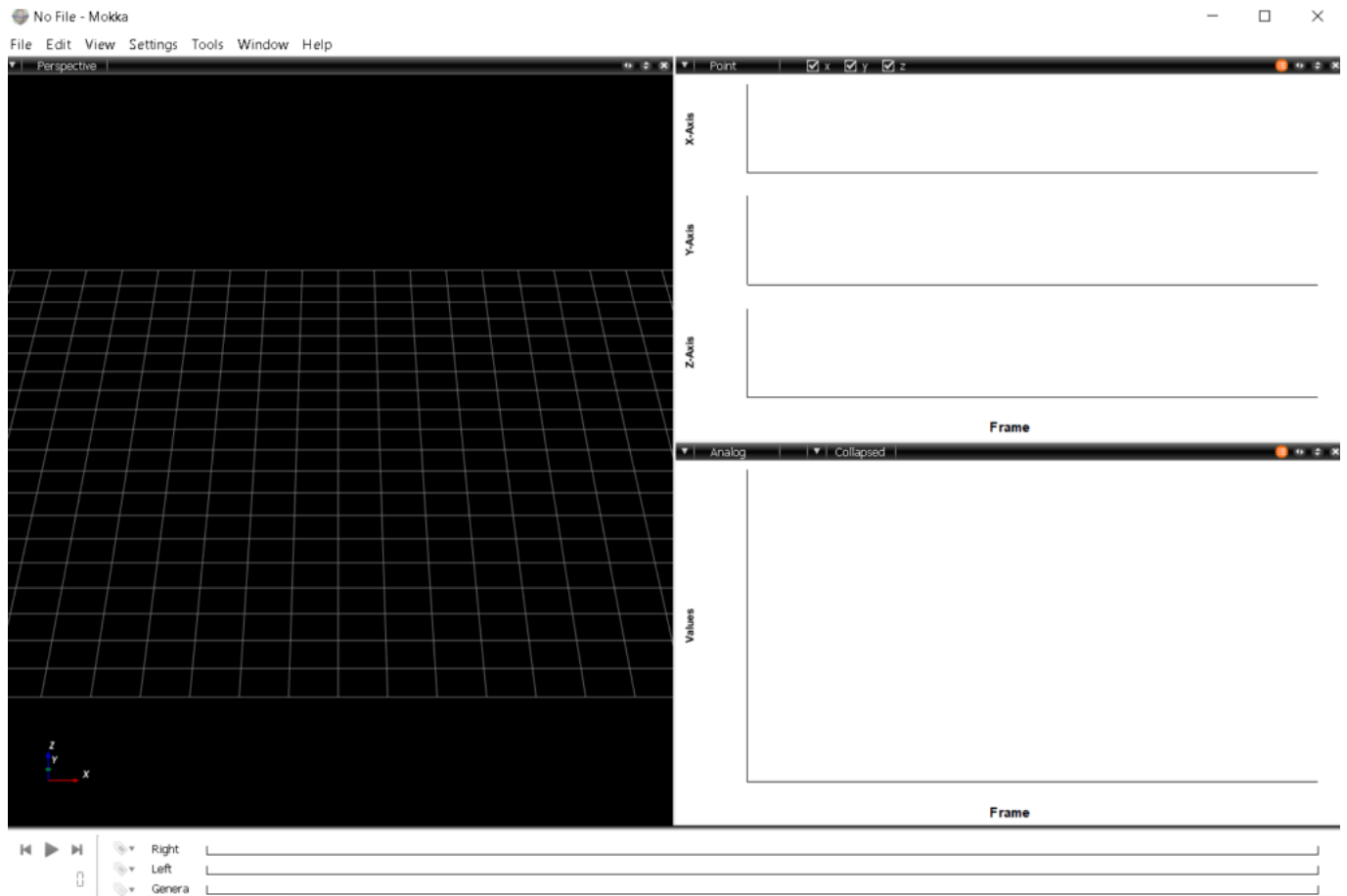


Figure 2: The default Mokka GUI

To load a C3D file, click on **File** and choose **Open**. Then you can choose a file that you wish to visualize. To try it out, you can download our [example data](#).

Please load *Running_Example.c3d* file. After that, the GUI should look similar to figure 3. Within the GUI you can explore the captured 3-dimensional experimental data. Each gray Point represents a spherical marker attached to the human body. By clicking on a specific marker, the name (often referred to as the marker label) is highlighted on the right-hand side **list box** (right side).

With the **play** button at the left lower corner of the GUI, you can see how the participant is running. Alternatively, you can use the **time bar** right next to the play button to inspect every frame (point in time) of your experimental data.



The **chart window** in the middle of the GUI can provide further information about the single components of your data. Note if the chart window is not shown go to **Window ->Layout -> 3D and Charts**. Now you can drag and drop labeled markers from the **List box** to the chart window, to see its X, Y, and Z coordinates over each instant in time.

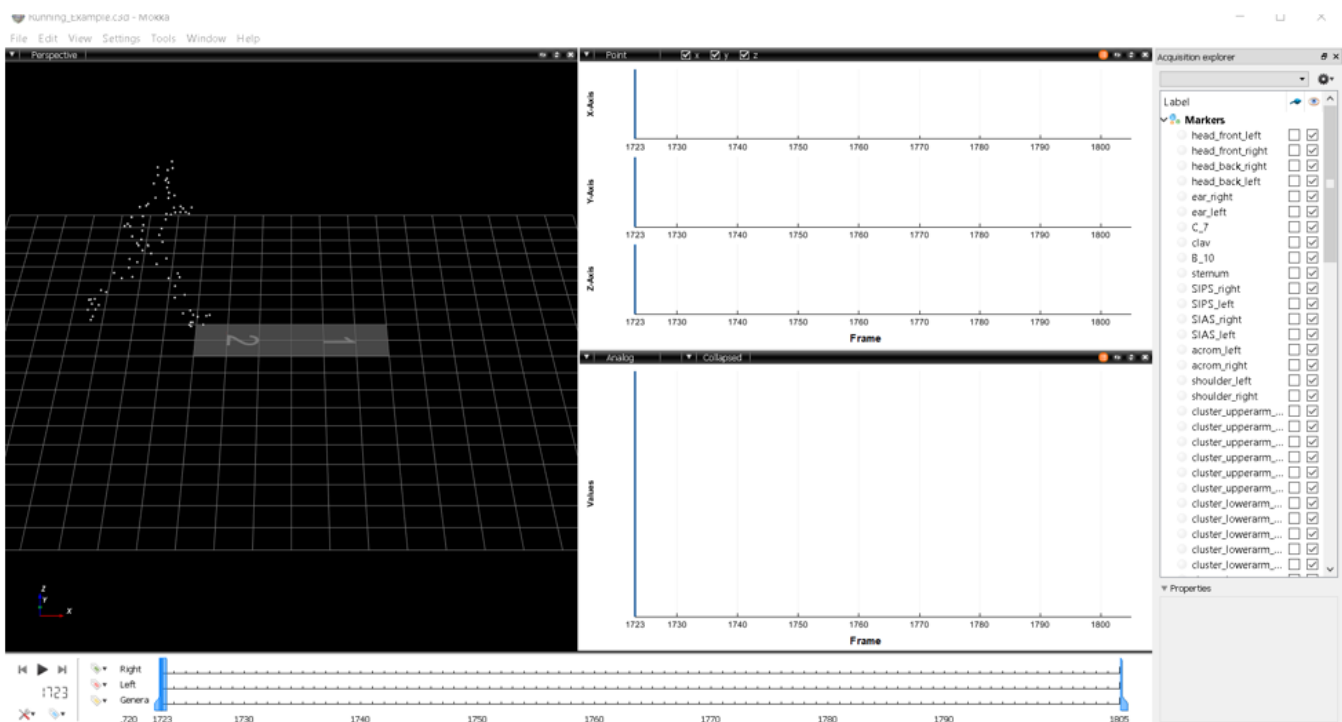


Figure 3: The Mokka GUI after loading Running_Example.c3d

Within the **list box**, you can also scroll down to the force platform data (if some exist). For the provided example Force Platform #2 was used. You can expand the force platform #2 field and can visualize the force components (Forces and Moments) in the same way as the marker trajectories by drag and drop the component to the chart view. However, when having many markers attached to your participant, visual inspection of your marker trajectories can become cumbersome. Therefore loading a model fitting to your data can be helpful.

For this tutorial, we have already generated a model for our marker setup. To load the model go to **Edit -> Options**. Check the box **Use default configuration** and click on **Choose ...** Select the file we have provided named **full_body_Model.mvc** . After the operation has



been completed the 3D view should contain different colored markers and connected segments.

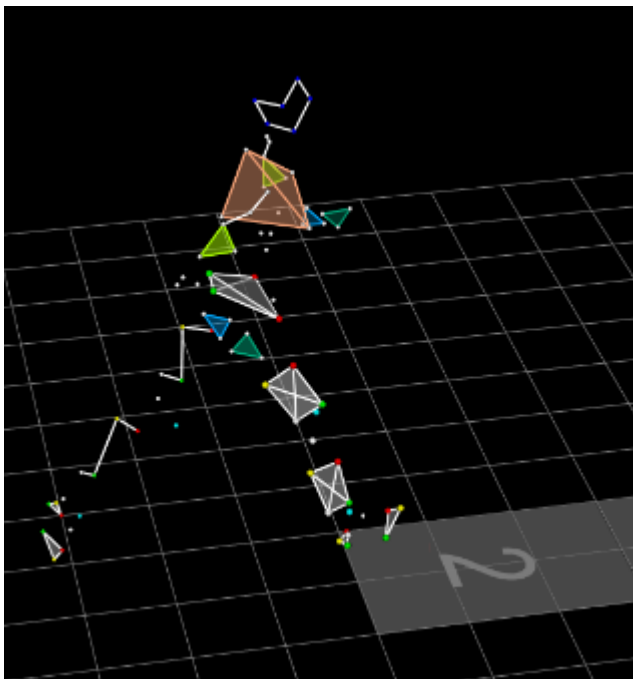


Figure 4: A whole body model adapted to our marker setup

Please note that the model has been adjusted to the used marker set. When using your own experimental data, with different marker names, you need to define new segments according to your marker labels. New Segments can be generated under **Tools -> Model -> New Segment**.

For further data processing, it is sometimes useful to crop your C3D files. Mokka provides the ability to crop (or cut) your C3D files very easily. Within the **time bar**, you can use the **sliders** to the left and the right end of the time bar to crop the C3D file. The frames which are removed from the C3D file are marked in blue. When right-click anywhere in the **time bar** you can select **Crop Frame of Interest**. Finally, you need to store the cropped file with **File -> Save As** or **Save** (to overwrite the existing file).

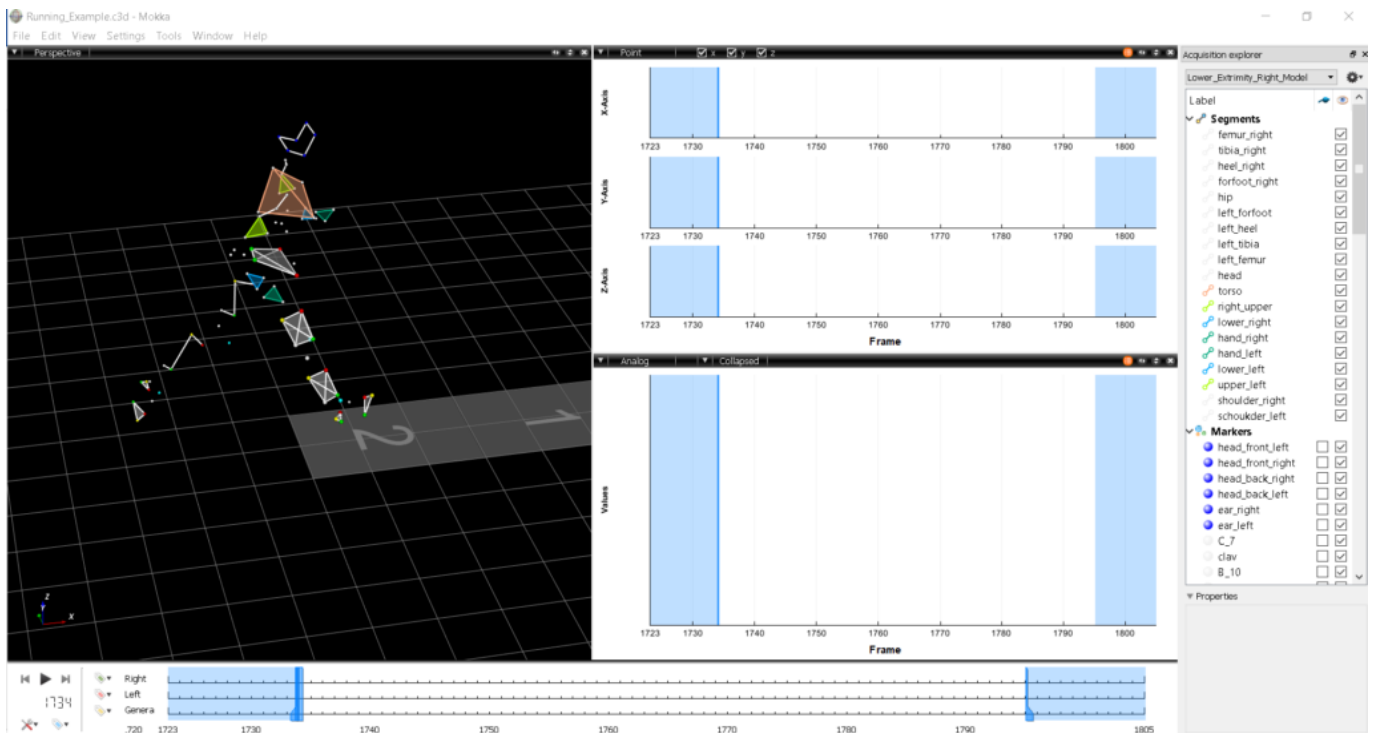


Figure 5: Cropping C3D files

Although, Mokka is useful for visualization purpose it does not allow to calculate joint kinematics and kinetics. It becomes also extraordinary time consuming when much experiential data needs to be batch processed. Within the next step, we would like to show you how to read in C3D files into Matlab.

2. Load C3D files with BTK in Matlab

For loading C3D files into your Matlab workspace you first need to download some additional, open-source functions. The **biomechanical toolkit (BTK)** is a frequently used cross-platform toolkit for reading, writing, and manipulating C3D files.

First, you need to download BTK from the website (<https://code.google.com/archive/p/b-tk/downloads>). Please ensure to download the correct version for your operating system. For us, the version [btk-0.3.0_Win7_MatlabR2009b_64bit.zip](#) works best. After downloading the toolkit you need to unpack the archive by using a standard archive converter.



Now you need to tell Matlab where all the BTK functions are located. Open Matlab and go to **Home Set Path Add with Subfolders...** and selected the folder (in our example `btk-0.3.0_Win7_MatlabR2009b_64bit`, note use the unpacked data folder and not the compressed zip archive). You can check if the Path is set correctly when typing `v = btkGetVersion` into your Matlab command window. If Matlab outputs a version number (for example '0.3.0') the path is set correctly.

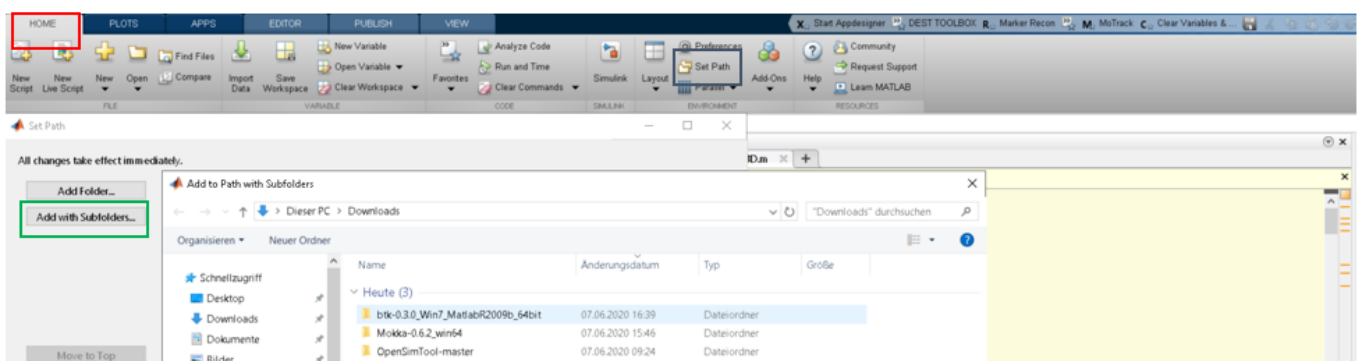


Figure 6: Ssetting the right folder path in Matlab

You can now read, manipulate, and write C3D files with Matlab. For the full list of the BTK function please refer to the following webpage

<http://biomechanical-toolkit.github.io/docs/Wrapping/Matlab/annotated.html>

In the [Download Section](#) below, you can find all example C3D data and some example Matlab code. This code will show you how to read and add virtual markers to a C3D file (static trial). Furthermore, the code will show you how to calculate stance time based on force data and add touchdown and toe-off events to a C3D file (running trial).

3. Downloads

- [Mokka](#)
- [Biomechanical toolkit \(b-tk\)](#)
- [Example C3D data and Matlab code](#)