

**Supplementary material:**

**A tutorial guide to ImageJ and Vessel\_width-plugin**

**Original article: Indirect measurement of the vascular endothelial glycocalyx layer thickness in human submucosal capillaries with a plug-in for ImageJ**

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## 1. Download ImageJ

Go to <http://rsbweb.nih.gov/ij/download.html> and download the version of ImageJ that matches the configuration of your hardware and operating system.



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### Download

#### Platform Independent

To install ImageJ 1.46 on a computer with Java pre-installed, or to upgrade to the latest full distribution (including macros, plugins and LUTs), download [ij146.zip](#) (3.2MB) and extract the ImageJ directory. Use the *Help>Update ImageJ* command to upgrade to newer versions.

#### Mac OS X

Download [ImageJ 1.46](#) (5.7MB) as a double-clickable Mac OS X application. Includes ImageJ64, which uses Java 1.6 in 64-bit mode on Intel Macs running OS X 10.5 or later. ([Instructions](#))

#### Linux

Download ImageJ 1.46 [bundled with 32-bit Java](#) (46MB) or [with 64-bit Java](#) (40MB). Both versions include Java 1.6.0\_24 from Sun and the ImageJ source code. ([Instructions](#))

#### Windows

Download ImageJ 1.46 [bundled with 32-bit Java 1.6.0\\_20](#) (28MB), [with 64-bit Java 1.6.0\\_20](#) (24MB; requires 64-bit Windows) or [without Java](#) (3MB). ([Instructions](#))

#### Documentation

Tiago Ferreira's comprehensive [ImageJ User Guide](#) is available as an 8MB PDF document and as a [ZIP archive](#). The [JavaDoc API documentation](#) is available as a ZIP archive.

#### Source Code

The ImageJ [Java source](#) consists of 92,000 lines of code in 265 files. It is available [online](#) and as a [zip archives](#).

#### Example Images

[31 downloadable sample images and stacks](#) are available in ImageJ's *File>Open Samples* submenu. These images, and more, are also available as a [8.2MB zip archive](#).

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You can also browse the ImageJ download directory at [imagej.nih.gov/ij/download/](http://imagej.nih.gov/ij/download/). Refer to the [release notes](#) for a list of new features and bug fixes.

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Save the ImageJ setup file on your computer and run the installation program. If necessary, use the ImageJ instructions on the download page ([Instructions](#)) to solve any platform specific problems encountered during the installation process.

## 2. Download the Vessel\_width.jar

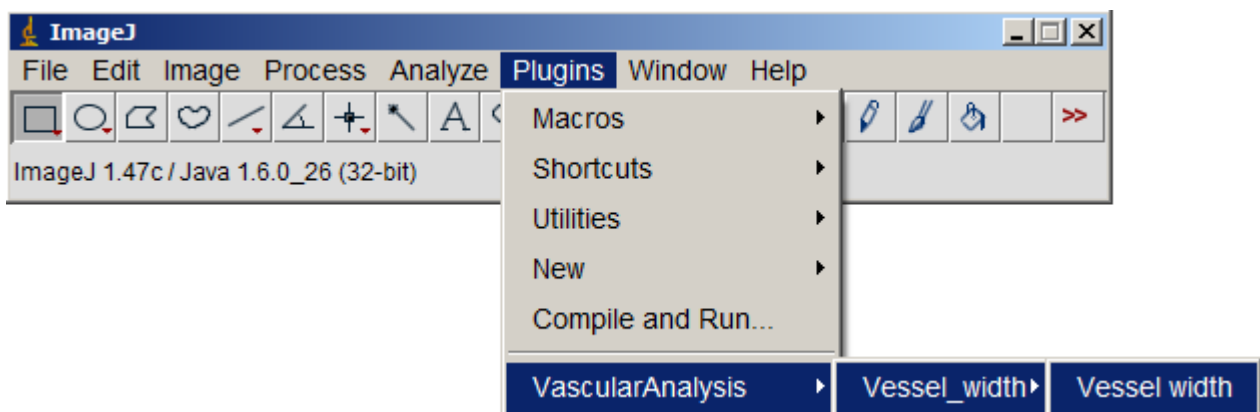
Go to <http://www.bioinformatics.org/microcirc/> and download the “Vessel\_width.jar” -file.

Place the “Vessel\_width.jar” file at the Plugins folder of the ImageJ directory.

## 3. Run ImageJ.

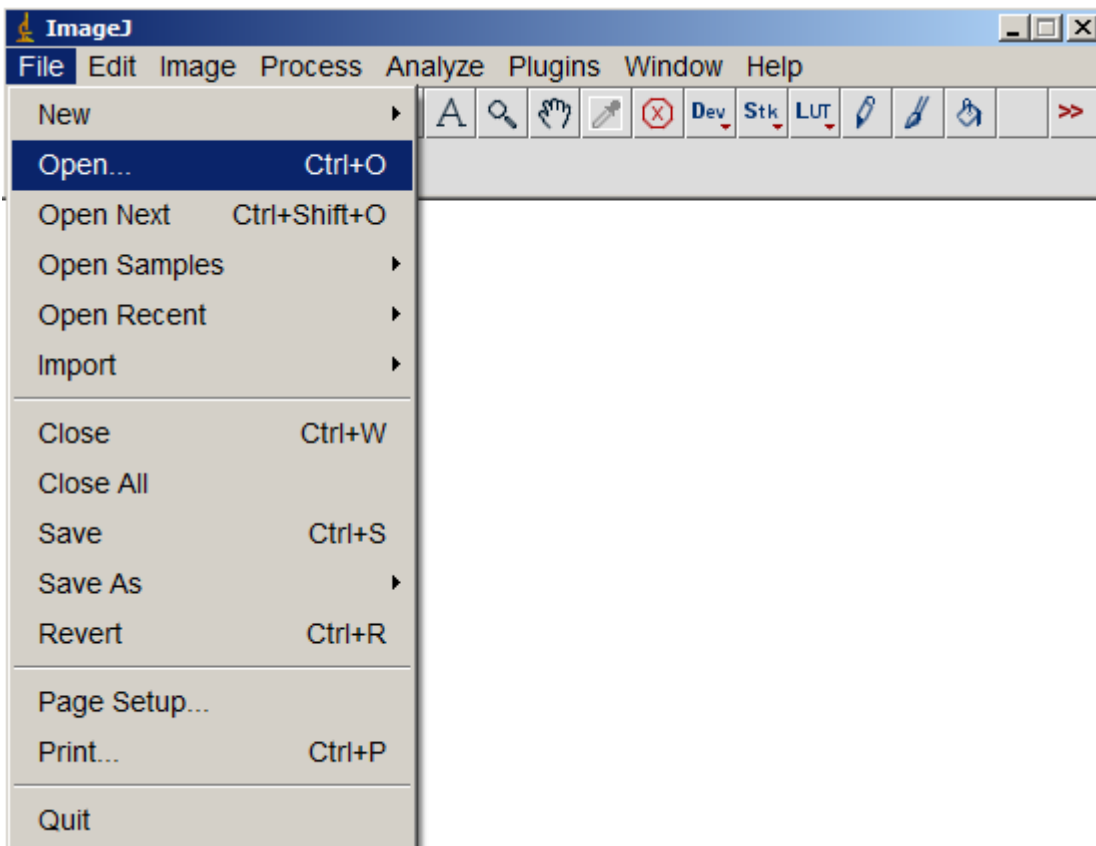
## 4. Locate the plugins icon and the Vessel\_width plug-in

The Vessel\_width plug-in should appear listed in the available plugins list under “Vascular analysis”. If not, make sure you have placed the Vessel\_width.jar in the correct folder of the ImageJ directory.

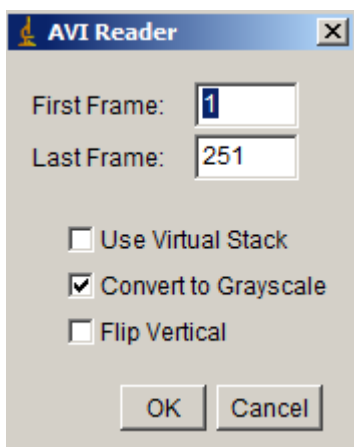


## 5. Open a microcirculatory AVI file with ImageJ

Open the AVI file by using the File -> Open command.

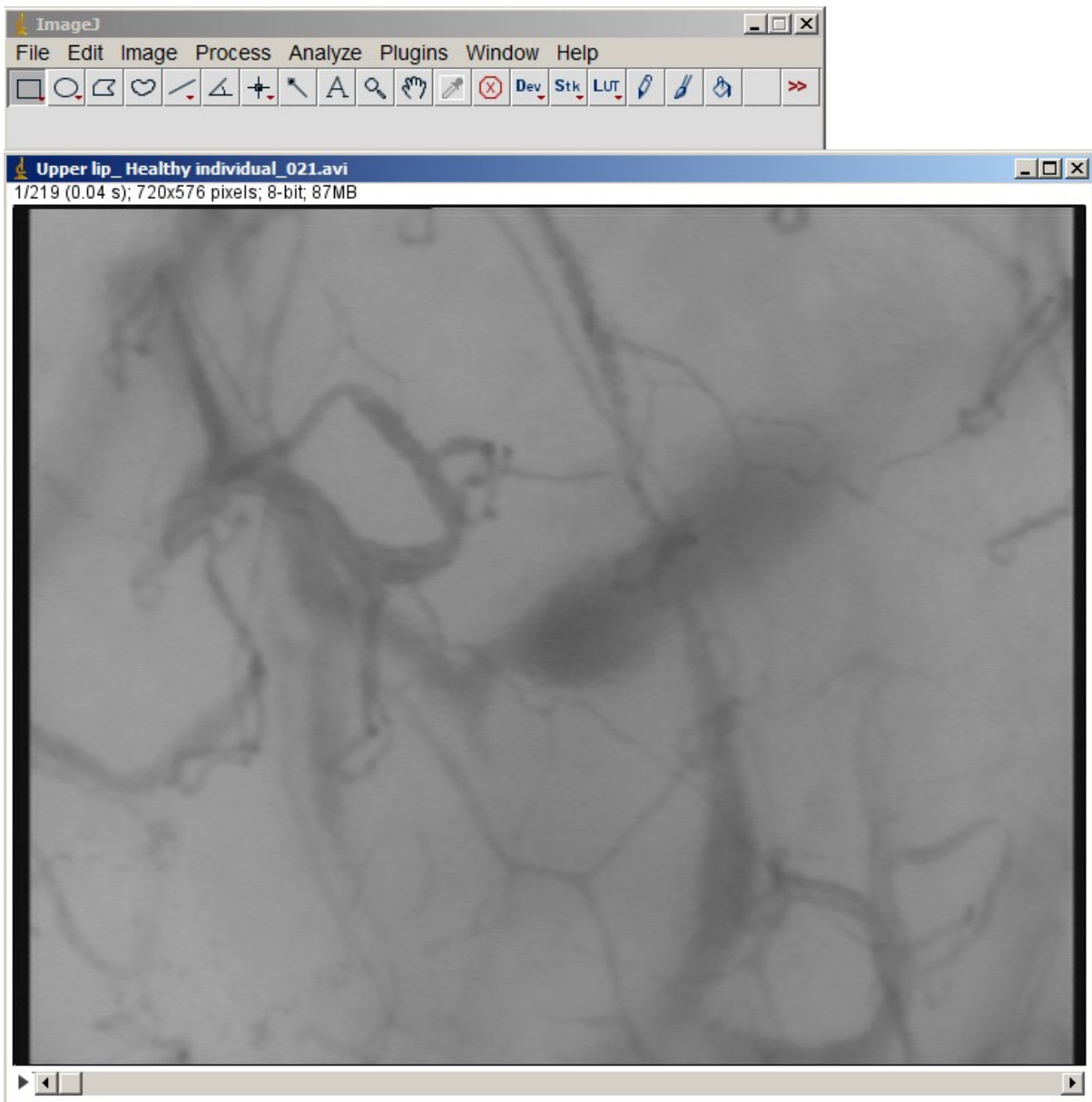


The ImageJ AVI reader box will appear. Choose the AVI frame numbers you want to open (default: First and Last frames of the file).



Optionally you may choose to convert the AVI file to 8-bit grayscale at this step. However, this is not necessary, as the Vessel\_width plugin will temporarily convert the opened AVI file to 8-bit grayscale to save memory and to make the measurement process faster.

An opened AVI file and the ImageJ command icons will appear like these:

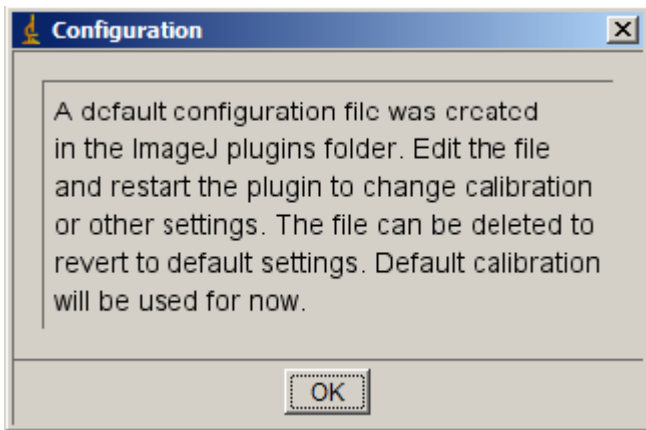


## 6. Activate the Vessel\_width.jar on top of the opened AVI file.

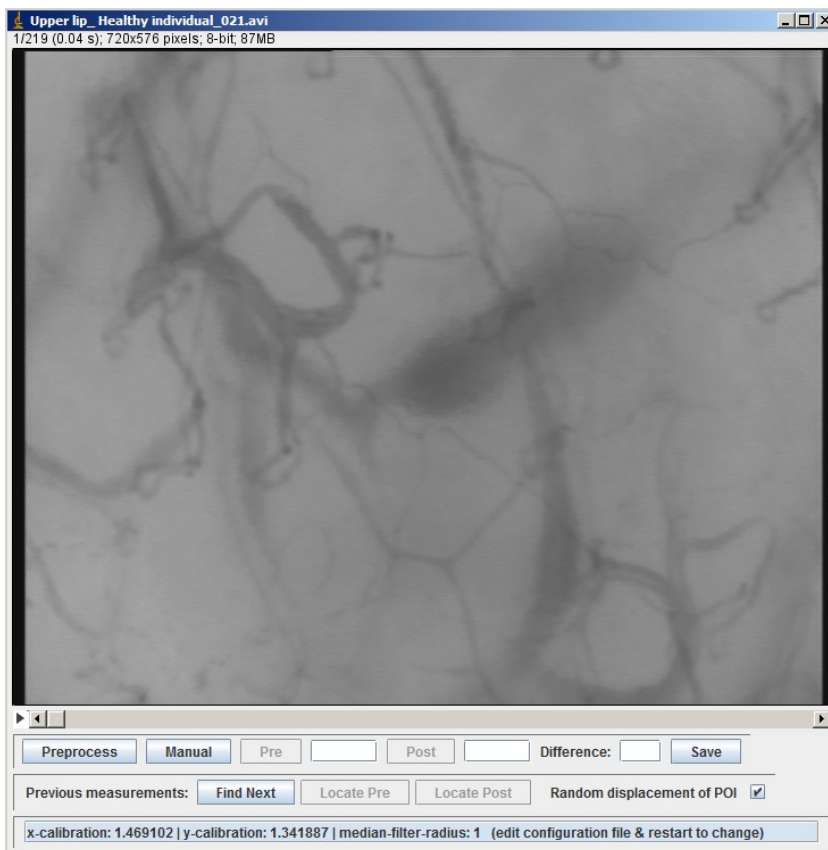
Follow the path: Plugins -> VascularAnalysis -> Vessel\_width ->Vessel\_width.

A default configuration file, (Vessel\_width-config.txt) will be automatically created in the plugins folder, when the Vessel\_width plug-in is used for the first time. The configuration (X- and Y-calibration, and the applied median filter radius) can be edited in a text editor, and saved to the Vessel\_width-config.txt -file in the plugins folder.

Press “OK”. (Exit ImageJ at this step to edit the configuration prior to any measurements, and restart).



The AVI-file will open with the Vessel\_width plug-in in a new window.



## 7. Locate a spontaneous white blood cell passage in a point-of-interest (POI) of a capillary.

Verify, that the shadow of the white blood cell also persists in the post-capillary venule (Figures A-D), before you start measuring the glycocalyx thickness.

**Figure A.** Point-of-interest in a capillary (beneath the asterix). The AVI frame preprocessed for demonstrative purposes (see the text of the original article for clarification of “preprocess”).



**Figure B.** A White blood cell (black-arrow) about to enter the capillary of interest. The AVI frame preprocessed for demonstrative purposes.



**Figure C.** A white blood cell passing the point-of-interest (black arrow). The AVI frame preprocessed for demonstrative purposes.



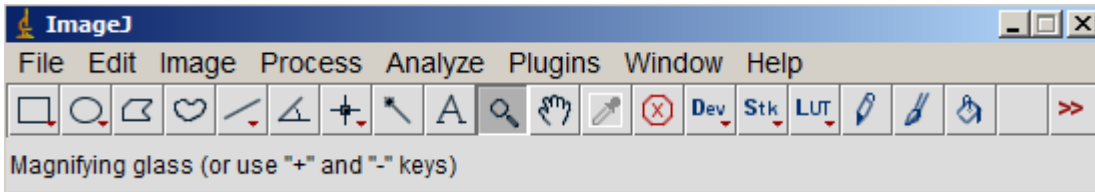
**Figure D.** The shadow of a passed white blood cell followed into a post-capillary venule (black arrow). The AVI frame preprocessed for demonstrative purposes.



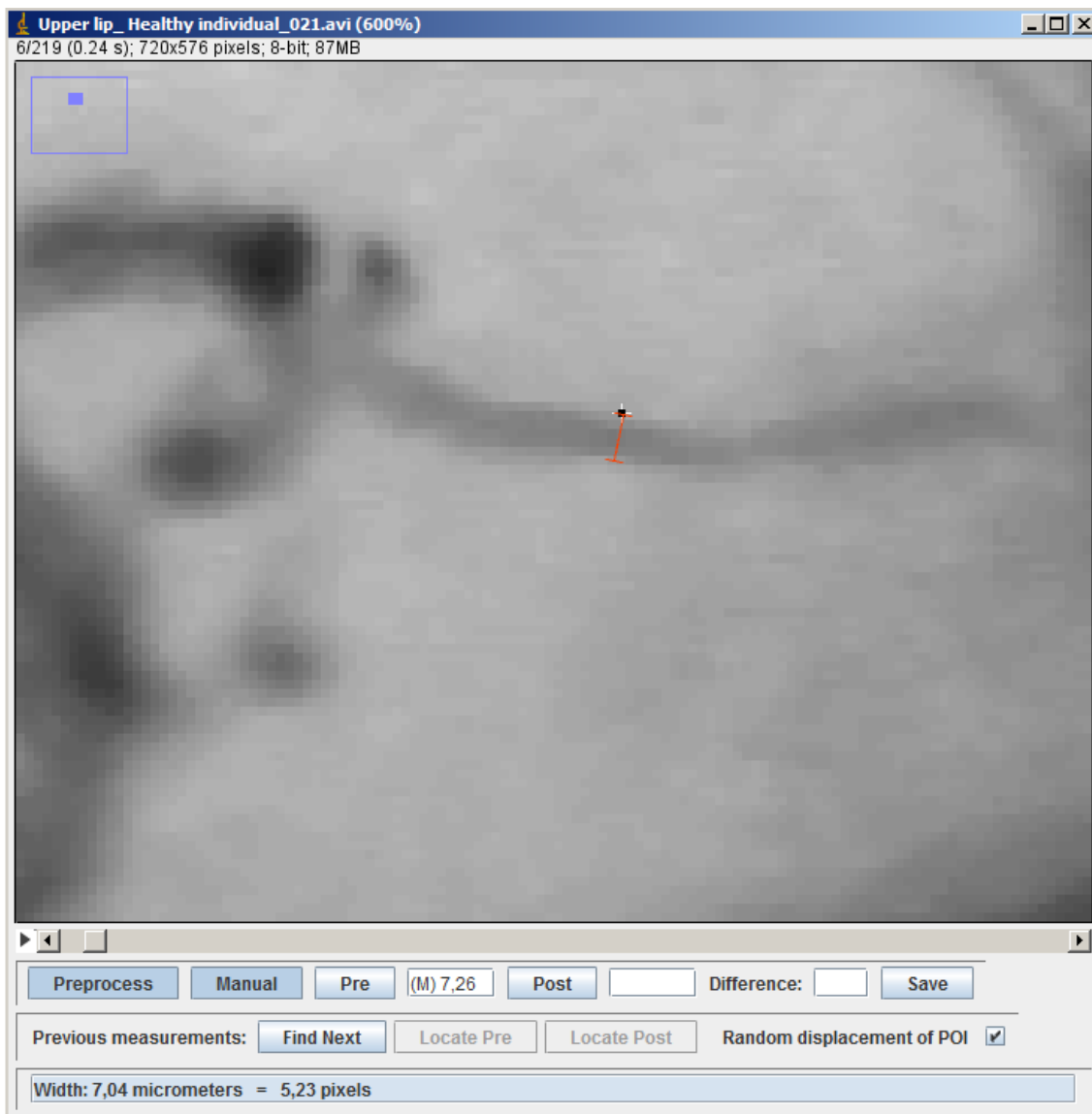


## 8. Manual measurements (Pre and Post).

Use the 600% magnification by activating the ImageJ magnifying glass tool on the image.



PRE: Activate the “Manual” icon of the Vessel\_width -plugin. Manual measurement tool (a free angle digital micrometer, red line on in the images below) will appear on the screen as you select a point on the image. A transection of the vessel (an ‘eye-ball’ estimate of diameter) can be drawn by drawing with a left mouse click. Visually approximate the diameter (edges of the red blood cell column) at the POI, where the WBC is about pass in the capillary. Then activate the “Pre”-icon, to measure the manually (M) drawn diameter.



POST: Measure the post diameter immediately after the passage of the WBC (need to scroll frames). Then activate the “Post”-icon, to measure the manually (M) drawn diameter.



Finally, press the “Save” icon to save the measurements in the log.

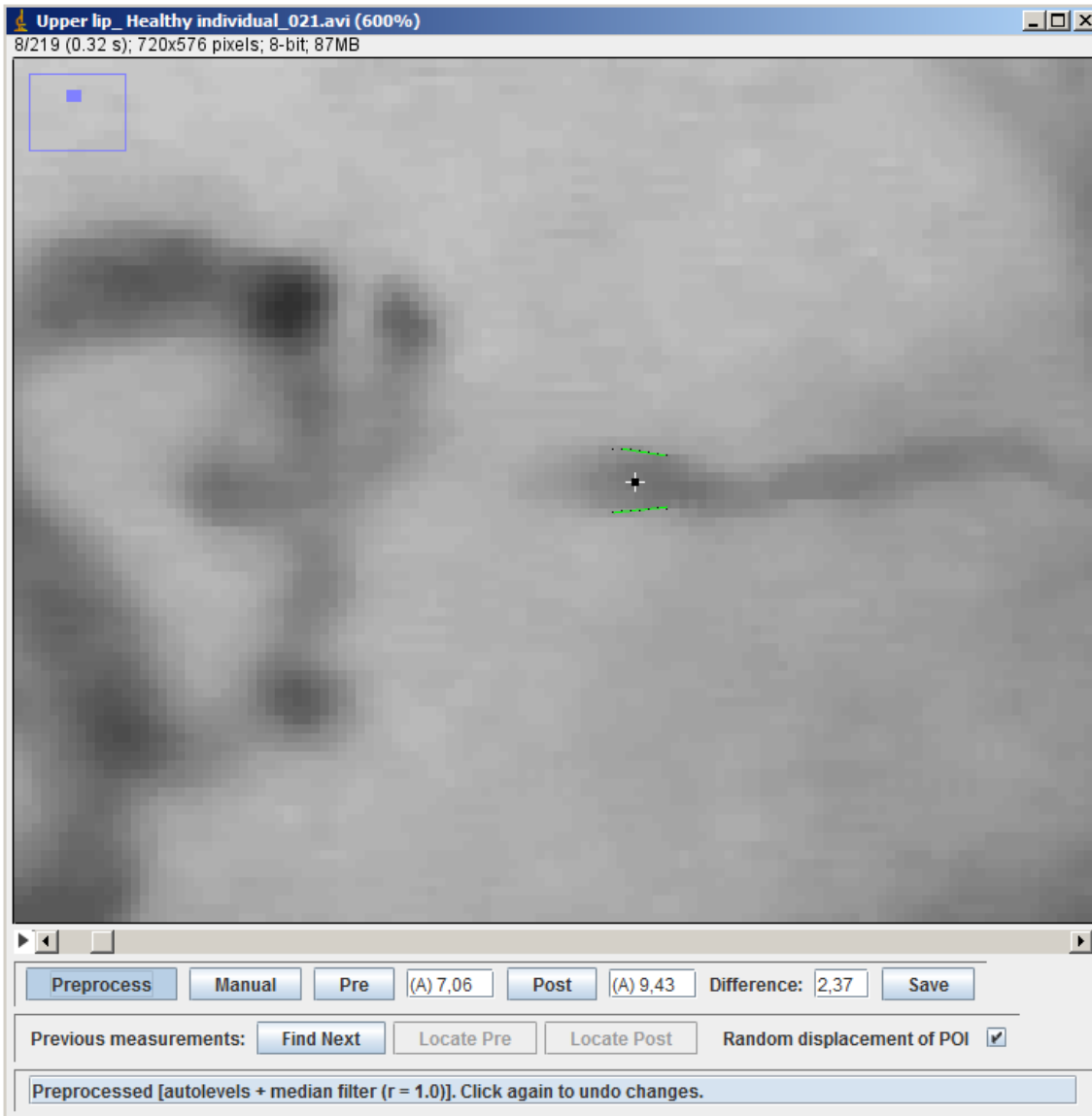
## 9. Semi-automatic measurements (Pre and Post)

Deactivate the “Manual” icon of the Vessel\_width -plugin (by clicking it), if it is active.

PRE: Place the point selection in the middle of the capillary at the POI, where the white blood cell is about to pass. Press the “Pre” icon, and the software will automatically detect the capillary thickness. Visually reassure the position and angle of detected red blood cell column edges (green lines). If the measurement is acceptable, proceed to post measurement.



POST: Place the point selection in the middle of the capillary at the POI, where the white blood cell has just passed. Press the “Post” icon, and the software will automatically derive the capillary thickness. Visually reassure the position and angle of detected red blood cell column edges (green lines).



Finally, press the “Save” icon to save the measurements in the log.

## **10. Troubleshooting:**

### **“No image - There are no images open”**

You attempt to run Vessel\_width plug-in before you have opened the AVI file. Press “OK” and proceed to open a microcirculatory AVI file first (See section 5).

### **Can't locate the Vessel\_width plug-in in the plugins tray**

Reposition the Vessel\_width.jar into the correct folder of the ImageJ directory.

You can even download the Vessel\_width.jar again, and place the reloaded file in the correct directory. Malpositioned .jar -files will not harm the software functions.