ActiPHV Manual

Installation

- (1) Install R (version 3.3.0) from: https://www.r-project.org/
 - a. If you already have R installed, but need to update the version, open R and type:
 - i. install.packages("installr")
 - ii. library(installr)
 - iii. updateR()
- (2) Install the required packages by typing:
 - i. install.packages("ggplot2")
 - ii. install.packages("reshape2")
 - iii. install.packages("FKF")
 - b. Check for any error messages.

Preparation

- (1) Copy the R-scripts **ActiPHV.R** and **startActiPHV.R** to your desired location and get the path (*scriptPath*) to this location.
- (2) Open startActiPHV.R in your editor of choice
- (3) Edit the parameters:
 - a. scriptPath = see (1). Important note: Make sure you use the right slash ("/"). If you copy the path from Windows Explorer, you will have to change the backslashes ("\") to slashes.
 - b. d = input directory. Should contain one file with tracking coordinates per filament (columns: Frame, x, y)
 - c. o = output directory. If it does not exist yet, it will be created. Important note: If you use the same output directory for different analyses, the results will be overwritten (not recommended). Please make sure not to create the output directory within the input directory.
 - d. f = frame rate in s/frame
 - e. $px = pixel scale in \mu m/px$
 - f. I = minimum phase length. All phases shorter than this will not be included into the results.

Start

- (1) Open R
- (2) Edit the parameters in startActiPHVV.R
- (3) Run startActiPHVV.R
- (4) The results will be in the directory you specified in the input parameters.

Results

- veloOfMaxFraction.txt: Summary for the maximum velocity result
- phaseResults.txt: All phase results with their corresponding filament IDs. For your analyses, the velocitiy/variance in μm/s (columns: "veloMicromPerSec"/"var_veloMicromPerSec") will be the

most important to analyze. If you want, you can check the visual phase results for every filament ID in the plot in *PhaseIdentification.pdf*

- PhaseIdentification2.pdf: Phase identification results without filament ID labels.
- DataForPlot.txt: Data used for the PhaseIdentification plots. Can be used to reproduce any plots, if necessary.