CAS3D DOCUMENTATION

V4.5 DOMINIK SCHNEIDEREIT, INSTITUTE OF MEDICAL BIOTECHNOLOGY

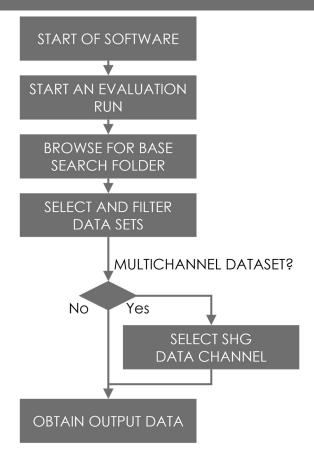
INTRODUCTION

This documentation offers an overview of the capabilities and the workflow of the CAS3D software package while also providing installation and setup instructions. A set of step-by-step instructions provides a detailed guideline for using the software.

INSTALLATION

Execute the installer package on Windows 10, 7 SP1 with a double click, and follow the installation instructions. If the Matlab 2019a runtime libraries are not already present on the machine, the installer will download and install the required files (Internet connection required). A startup shortcut is placed on the desktop if the respective option is selected, and the default installation folder is "C:\Program Files\Institute of Medical Biotechnology\CAS3dTool". The software can be started by double-clicking the desktop shortcut or the CAS3dTool.exe (the default location of the executable file is: C:\Program Files\Institute of Medical Biotechnology\CAS3dTool\application).

USAGE FLOWCHART



USAGE STEP-BY-STEP

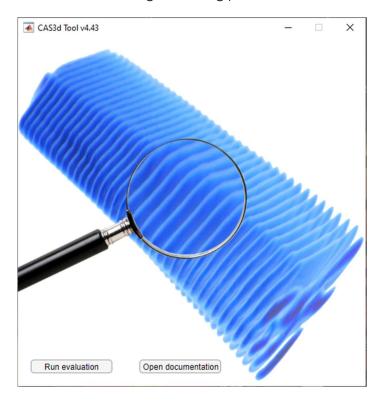
START OF SOFTWARE

The software is launched by double-clicking the desktop icon. After a short loading time that depends on system specifications, a console window is displayed, which will provide basic information and feedback while the software is running, initially displaying the software version. Closing the console window will exit the software, terminating all running processes.



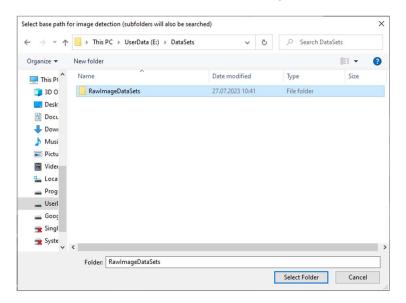
THE LAUNCHER WINDOW

The launcher window appears alongside the console window shortly after starting the software. An evaluation run can be started and this documentation can be accessed from the launcher window. If the launcher window is kept upen, multiple evaluation runs may be performed in sequence without the need to re-load the Matlab runtime engine. Closing the launcher window will exit the software, terminating all running processes.



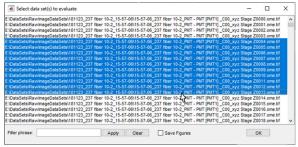
BROWSE FOR BASE SEARCH FOLDER

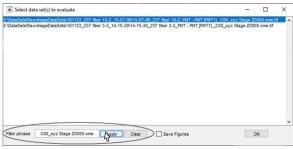
A file browsing dialogue is displayed where a folder is to be selected. The folder marks the base search folder, the whole folder tree within the marked folder or drive will be scanned for data files in the .ome.tiff, tiff or .tif format, that will be listed in the next step. The dialogue will open in the folder location that was last indicated for an evaluation. When the folder does no longer exist, or the software is restarted, the folder to start browsing from will default to "C:\".



SELECT AND FILTER DATA SETS

The list of detected files is displayed, showing their full path to allow for unambiguous identification of each file. One or multiple files may be selected for evaluation by a combination of the common drag-click, shift-click, and control-click commands that are applicable in most file browser dialogues. Additionally, a filter string may be specified in the lower left of the panel that, when applied, will show a reduced list of files that contain the specified string in their name or path. The previously mentioned multi-selection methods may still be applied on the filtered file list. An applied filter string will be retained and applied automatically when the software is executed the next time but can be changed or cleared if desired. Be sure to only select one slice per image stack, if the image slices of your stacks are saved as individual files. Also be sure the selected slice contains the stack metadata (usually this is the first slice of a stack). Otherwhise, the evaluation will fail or repeat the stack evaluation for every selected slice. The current selection is confirmed, and the software will proceed to the next dialogue by a click on OK. If the "save figures" checkbox option is selected a summary figure is created and saved for each evaluated data set. Unchecking this option will skip the generation of the output figures and reduce the required runtime for an evaluation run.

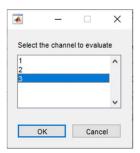




UNFILTERED FILTERED

SELECT CHANNEL

If multiple channels are detected in the selected data set, the software will prompt the user to pick the channel that contains the SHG signal to be used for evaluation. Other channel data are not relevant in the evaluation and will be ignored. Confirm the relevant channel by double clicking or selecting it and clicking the OK button.



ANALYSIS PROGRESS

During the analysis process, the progress through the data sets is displayed in a progress bar. Intermediate results and prorgress steps are displayed in the console window. Warnings and error messages are also displayed in the console window. An output graph for each data set are displayed after a successful evaluation.

```
Saturation: 1% Background: 78.9519%
Finding dominant pattern and its orientation through iterative Fourier transform analysis

Iterations neccessary to find secondary peaks: 6
Detected main orientation 104.1451 ° and elevation 10.4729 ° with mean SL: 2.6995 micron

CAS3D = 0.80832 wCAS3D = 0.82601
Processing: 181123_237 fiber 3-3_14-15-30_14-15-30_237 fiber 3-3_PMT - PMT [PMT1] _C00_xyz Stage Z0000.ome

Reading series #1

Analyzing Images:1/2

Saturation: 1.0001% Background: 80.9164%
Finding dominant pattern and its orientation through iterative Fourier transform analysis

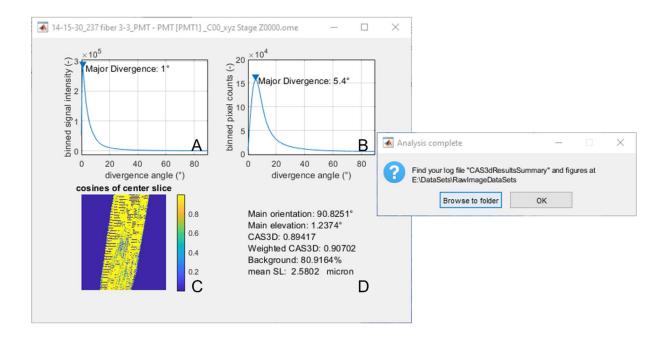
Iterations neccessary to find secondary peaks: 6
Detected main orientation 90.8251 ° and elevation 1.2374 ° with mean SL: 2.5802 micron

CAS3D = 0.89417 wCAS3D = 0.90702

Analysis complete!
Find your log file "CAS3dResultsSummary" and figures at E:\DataSets\RawImageDataSets
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ANALYSIS RESULTS AND OUTPUTS

After the evaluation run is complete and all data sets have been processed. A dialogue window informs of the successful completion and of the number of warnings during execution. The dialogue also indicates the location of the results log file. When clicking "Browse to folder", a windows file browser window is opened at the location of the log file. The log file is a spreadsheet containing all evaluated parameters of all data sets. An output graph summarizes the results of each analyzed data set. The output graphs are saved alongside the log file in .png and Matlab .fig format. The summary graphs shows a histogram of voxel gradient orientation divergence unweighted (A) and weighted by voxel intensity (B). It shows the center image slice of the data set with the cosine angle of each voxel is shown in a false color scale. Additionally, the key output parameters of the evaluation are listed in the output graph and the file name of the data set is displayed in the window header.



LOG FILE STRUCTURE

The log file, named "CAS3DResultsSummary" is provided as a basic spreadsheet in commaseparated values format(.csv). The separator character is defined by your windows system settings. Depending on your region settings, the separator may be a different character, such as a semicolon or a tabstop. CSV files are compatible with most spreadsheet software such as Excel, OpenOffice Calc or Google Tables. Each measurement run creates a new log file. In case of duplicate file names, an incrementing four-digit number is appended to the end of the file name.

The log file contains a simple data structure in which one line represents the results of a single volumetric image, and each column represents a descriptor or output value. The Filename column contains the full folder path that points to the evaluated volumetric image and serves as a unique identifier. The evaluated parameters CAS3D, intensity weighed CAS3D (intWeighedCAS3D), dominant Orientation Angle (MainOrientation), dominant elevation angle (MainElevation), mean angular deviation (AngularDeviation), mean intensity weighed angular deviation (WeighedDeviation), and mean sarcomere length (Mean SL) are provided as fixed point values with four digit precision. The image saturation (Saturation) and image backround (Background) are provided as floating point values with four digit precision.

