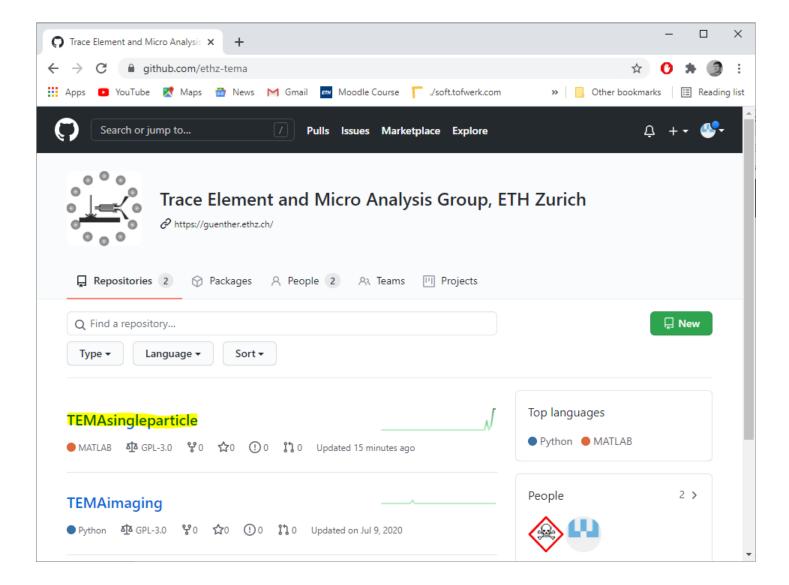
Step by step NanoFinder instruction

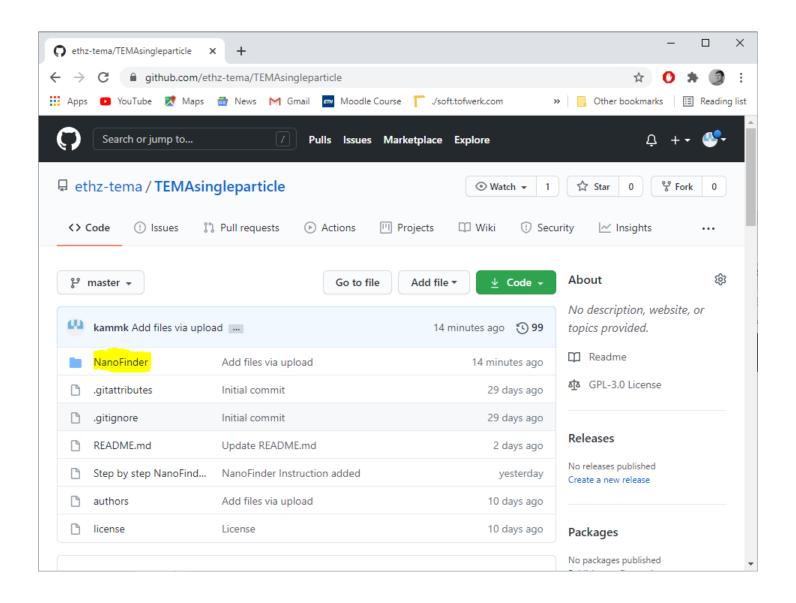
Author: Kamyar Mehrabi: kamyarm@ethz.ch

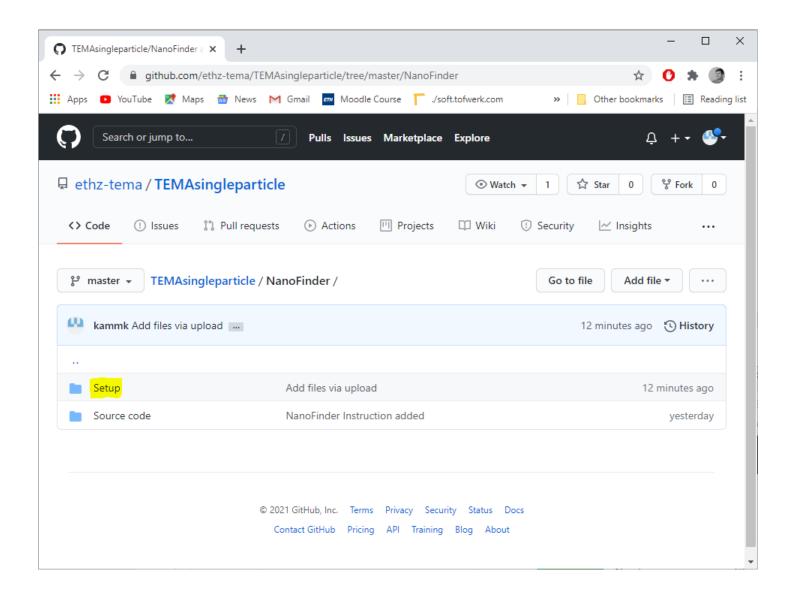
Installation:

1. In order to download, please go to the link below and download the contents of Setup folder

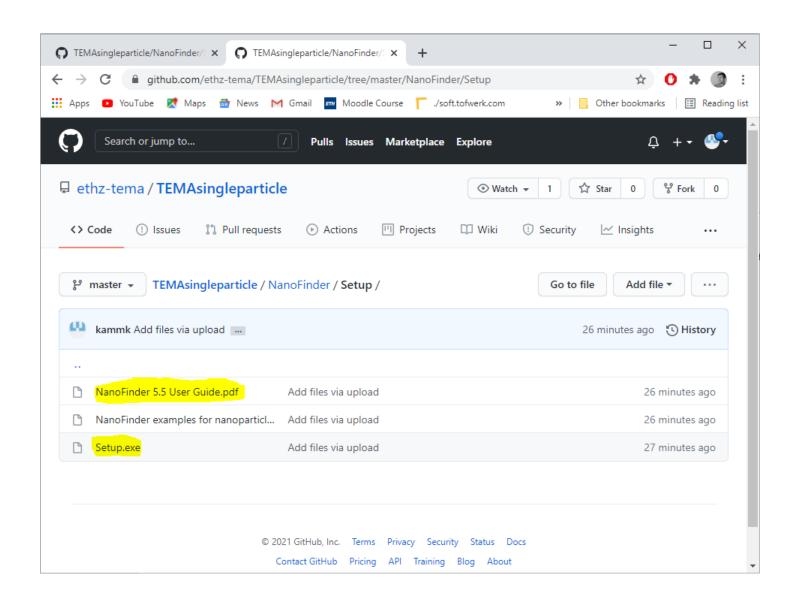
Link: https://github.com/ethz-tema

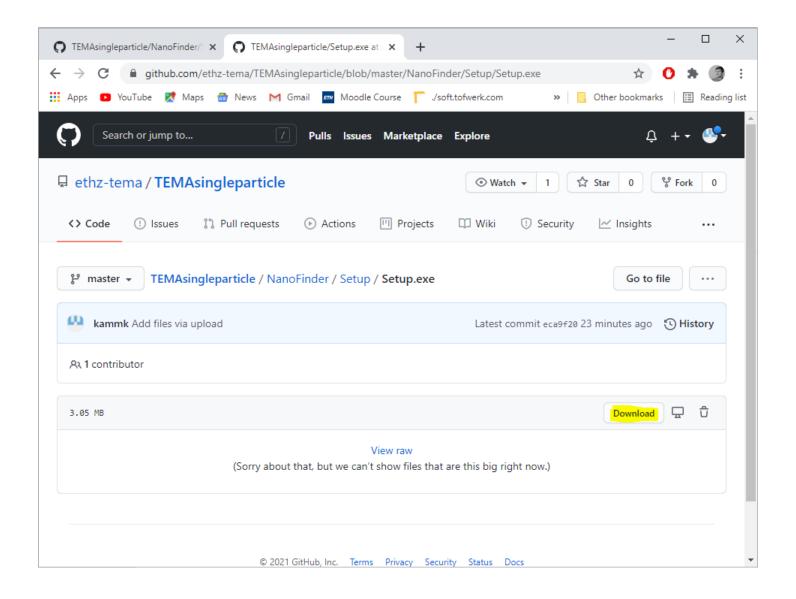




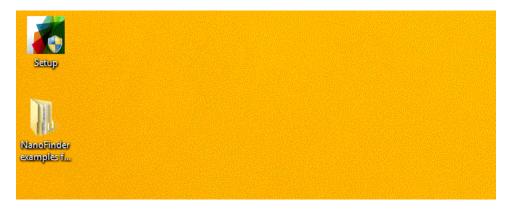


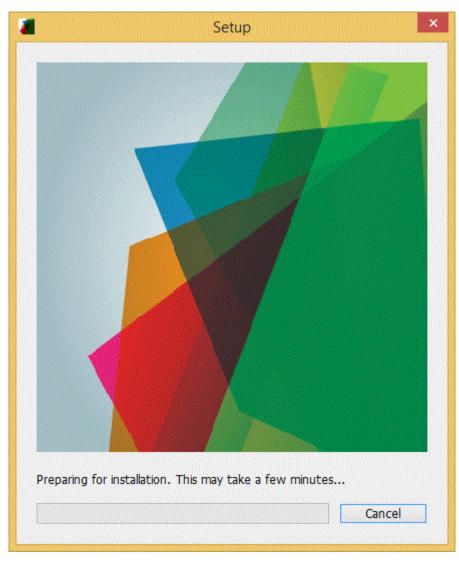
2. Download the **Setp.exe** and **NanoFinder User guide.pdf** from Setup directory as shown in the following screenshots.

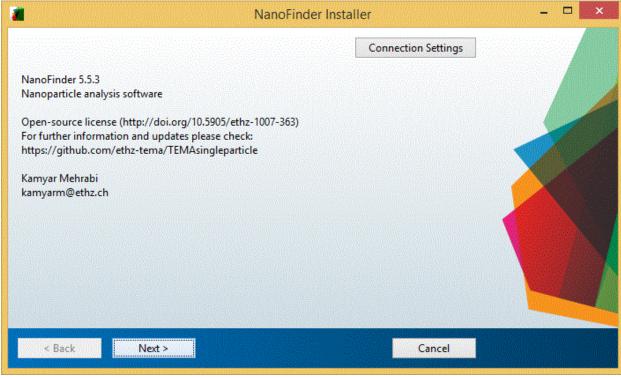


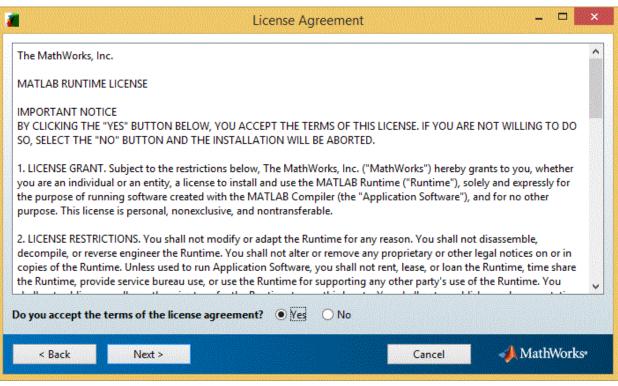


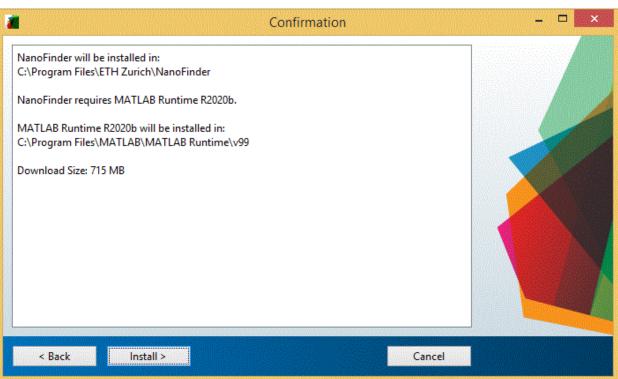
3. Run the **Setup.exe** and follow the steps as being shown in the pictures below.

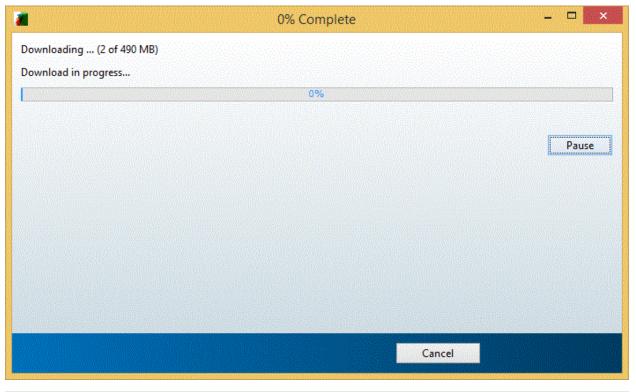


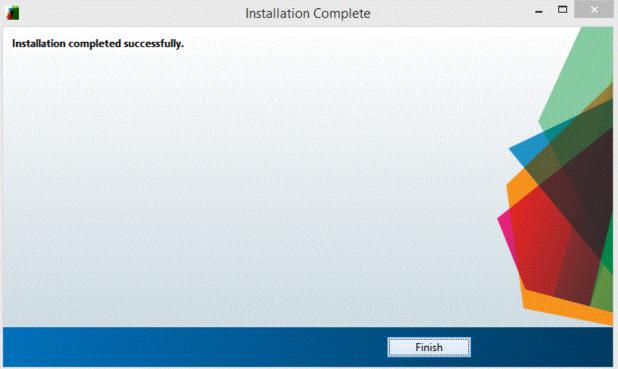












4. After installation is finished, you could find the NaFinder program in your programs directory. Search for NanoFinder in your programs directory and copy a shortcut of it in Desktop or location of your choice.



Processing the test sample:

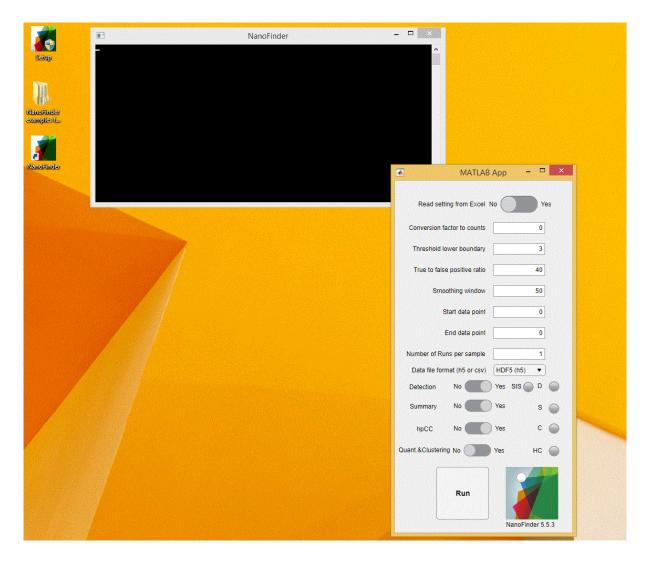
Complete example data files could be obtained from link below. After download please unzip the folder.

https://polybox.ethz.ch/index.php/s/5GeuULpyiBwBIDT

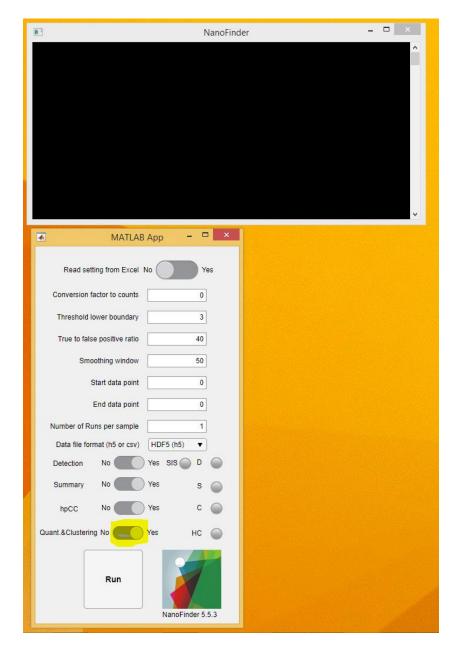
- 1. Open the NanoFinder program
- 2. A terminal window will show up shortly after



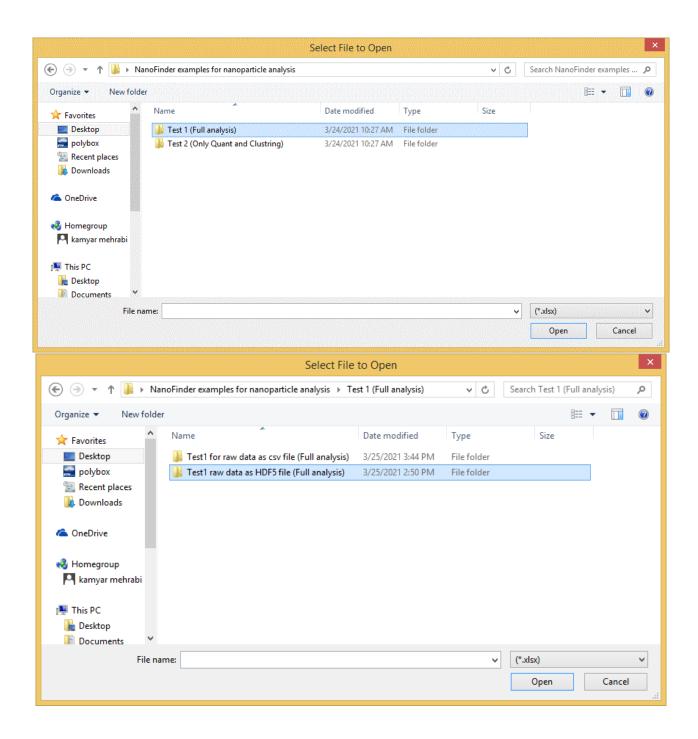
3. Wait until the program graphical user interface (GUI) open. It will take a few seconds to minutes depending on the processing power of your computer.



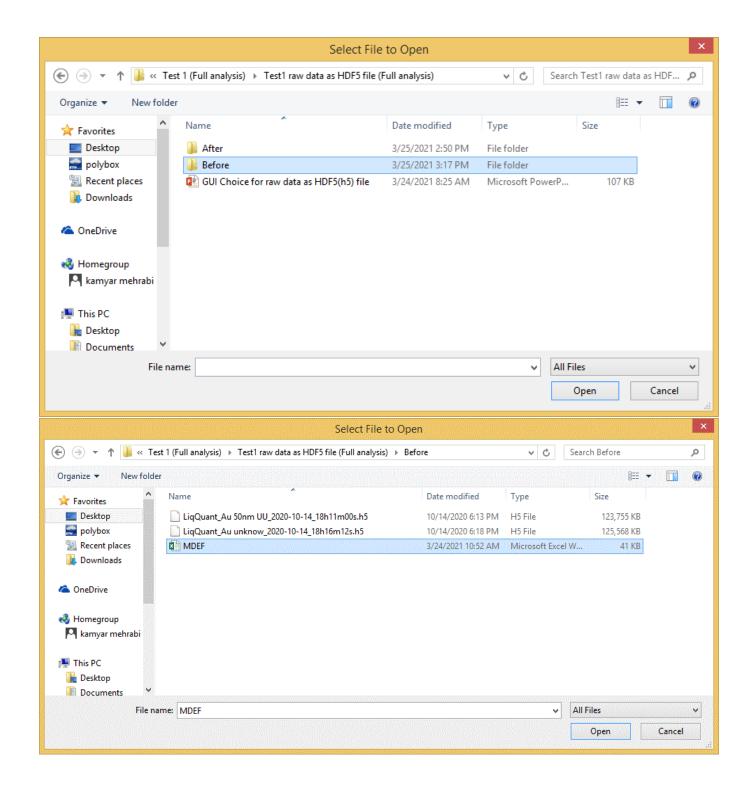
4. Fill in the GUI as being shown below (also in the NanoFinder test folder PowerPoint file). Here we processed the **Test 1 HDF5 file and read setting from GUI**, so the GUI will look like the following after we fill it correctly



5. After you fill the GUI, press Run. Then a file browser will popup. direct the file browser to the location of your raw data directory, where you also copied the MDEF.xlsx.



6. Here we want to process the **Test1 raw data as HDF5 file**. In that folder go to the **Before** file and **open**MDEF.xlsx



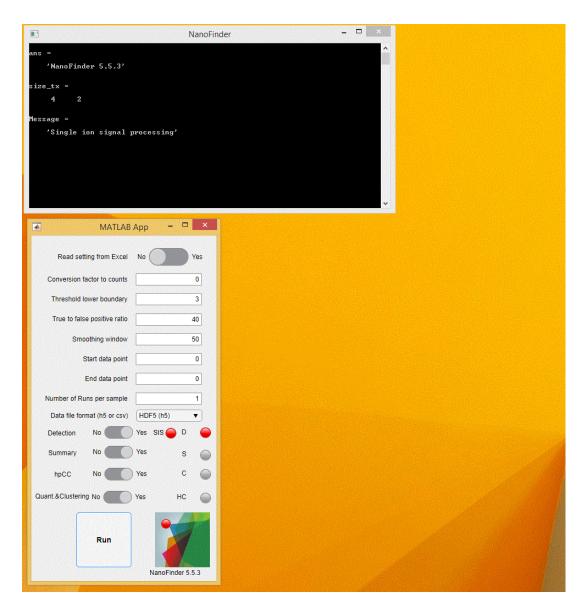
7. Program start processing as following screenshots is showing.

The terminal window, indicator lights in GUI, and popup progress charts will inform you about the latest analysis that is being done.

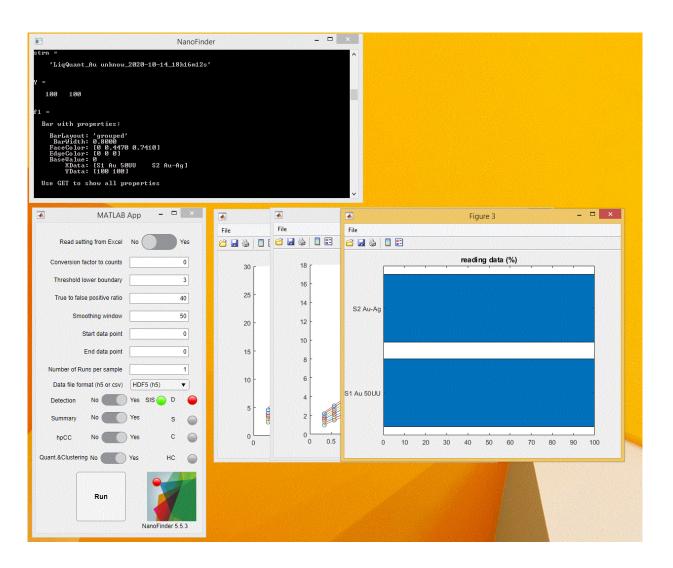
*While the program is running, it is highly recommended to prevent using Excel, since the program is using it.

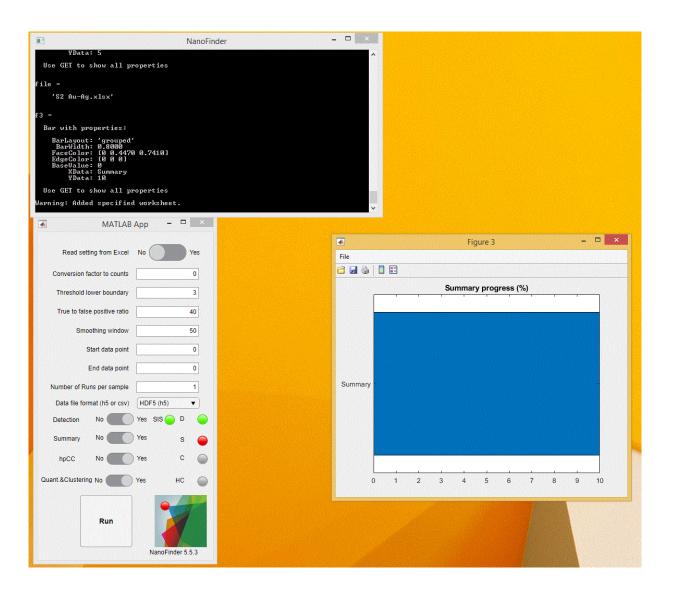
In case any error happens, see the terminal window and read about the error there carefully. In case the error is not solved, an Email may be sent with **Nanofinder run status** file attached to it. The Naofinder run status is generated as soon as you open the Nanofinder program and it is saved at the location of your running Nanofinder program shortcut. It contains the same data as shown in the terminal window.

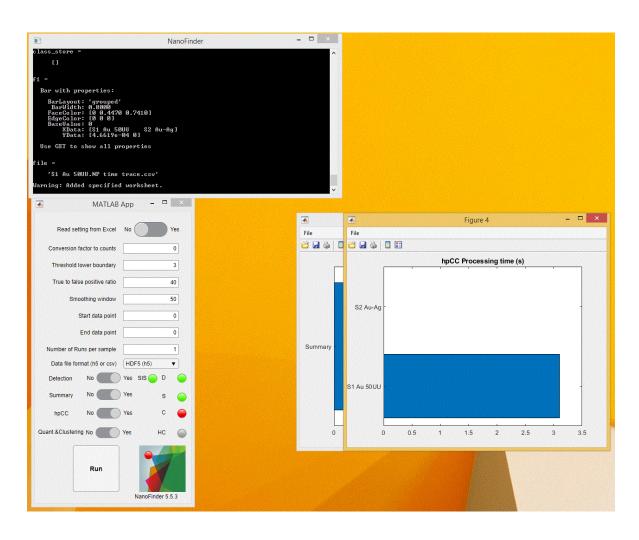


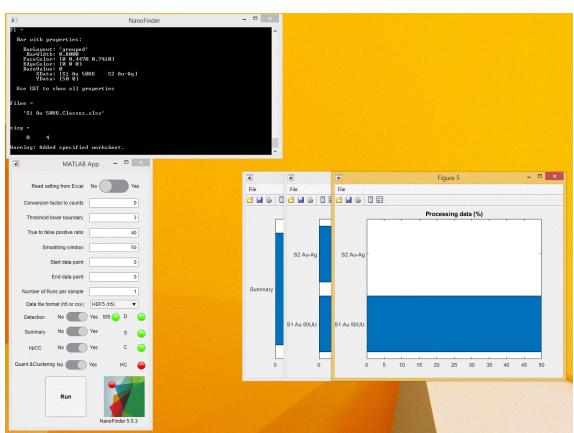


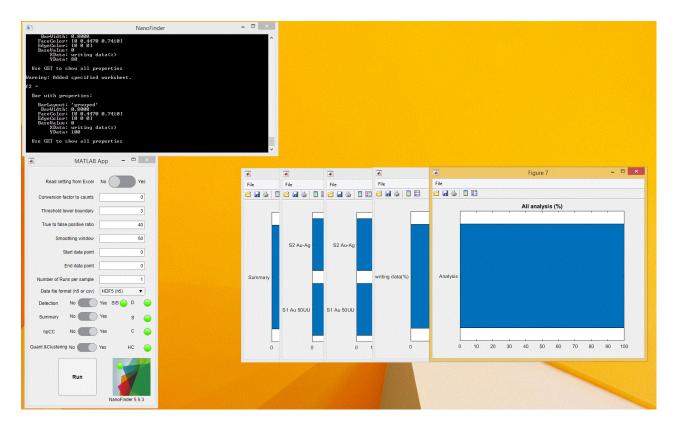




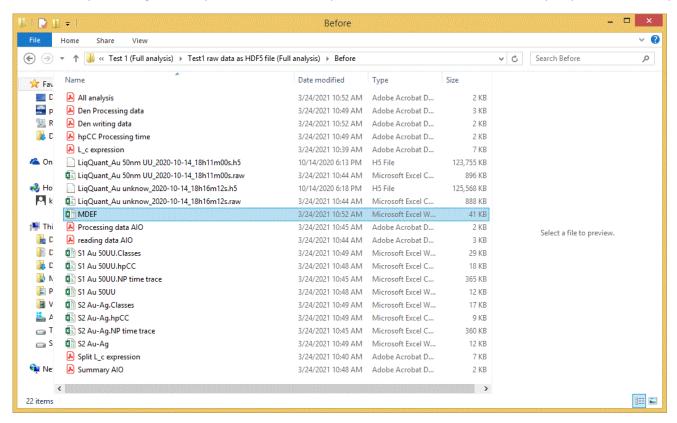








8. After all processing is done, you could access the processed data in the same directory as your raw data input.



For more information and a guide on how to set up your analysis please go to the GitHub folder and find the Nanofinder user guide.pdf in the same directory as Setup.exe.

- 1. K. Mehrabi, R. Kaegi, D. Gunther and A. Gundlach-Graham, Emerging investigator series: Automated Single-Nanoparticle Quantification and Classification: A Holistic Study of Particles into and out of Wastewater Treatment Plants in Switzerland, *Environ. Sci.: Nano*, 2021, DOI: 10.1039/D0EN01066A.
- 2. A. Gundlach-Graham, L. Hendriks, K. Mehrabi and D. Gunther, Monte Carlo Simulation of Low-Count Signals in Time-of-Flight Mass Spectrometry and Its Application to Single-Particle Detection, *Anal. Chem.*, 2018, **90**, 11847-11855.
- 3. A. Gundlach-Graham and K. Mehrabi, Monodisperse microdroplets: a tool that advances single-particle ICP-MS measurements, *J. Anal. At. Spectrom.*, 2020, **35**, 1727-1739.
- 4. K. Mehrabi, D. Gunther and A. Gundlach-Graham, Single-particle ICP-TOFMS with online microdroplet calibration for the simultaneous quantification of diverse nanoparticles in complex matrices, *Environmental Science-Nano*, 2019, **6**, 3349-3358.