

This vascular analysis runs inside of [Igor Pro](#).

Open bHyperstack in Igor Pro

- Double-click on desktop icon bHyperstack.
- Inside Igor, activate the Hyperstack menu by clicking on the Igor command window. The Igor command windows title is 'Untitled'.
- Select menu 'Hyperstack -> Load User File' and select your .txt file.

Load a hyperstack

- In main [Hyperstack panel](#), click 'Load' button.
- You then load a hyperstack by opening a file ending in _T.ibw
- For example if your hyperstack is named a152hs1, *you would load the file 'a152 hs1_T.ibw'*

Save a hyperstack

- Saving is done one hyperstack at a time.
- Select a hyperstack in main [Hyperstack panel](#) and click 'Save' button. The hyperstack will be saved on the hard-drive in the folder it was loaded from.

Quit Igor

- Select menu 'Igor Pro -> Quit Igor Pro
- You will be asked 'Do you want to save changes to experiment "Untitled" before quitting'. Answer No.

Important: You are responsible for saving your work for each hyperstack. When you quit Igor Pro, your work is NOT AUTOMATICALLY SAVED.

You save your hyperstack using the save button in the main [hyperstack panel](#).