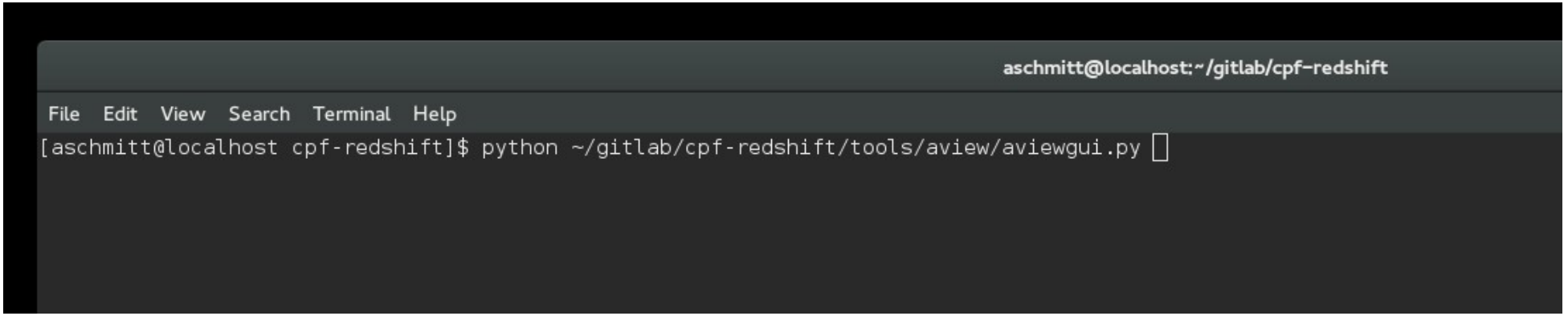
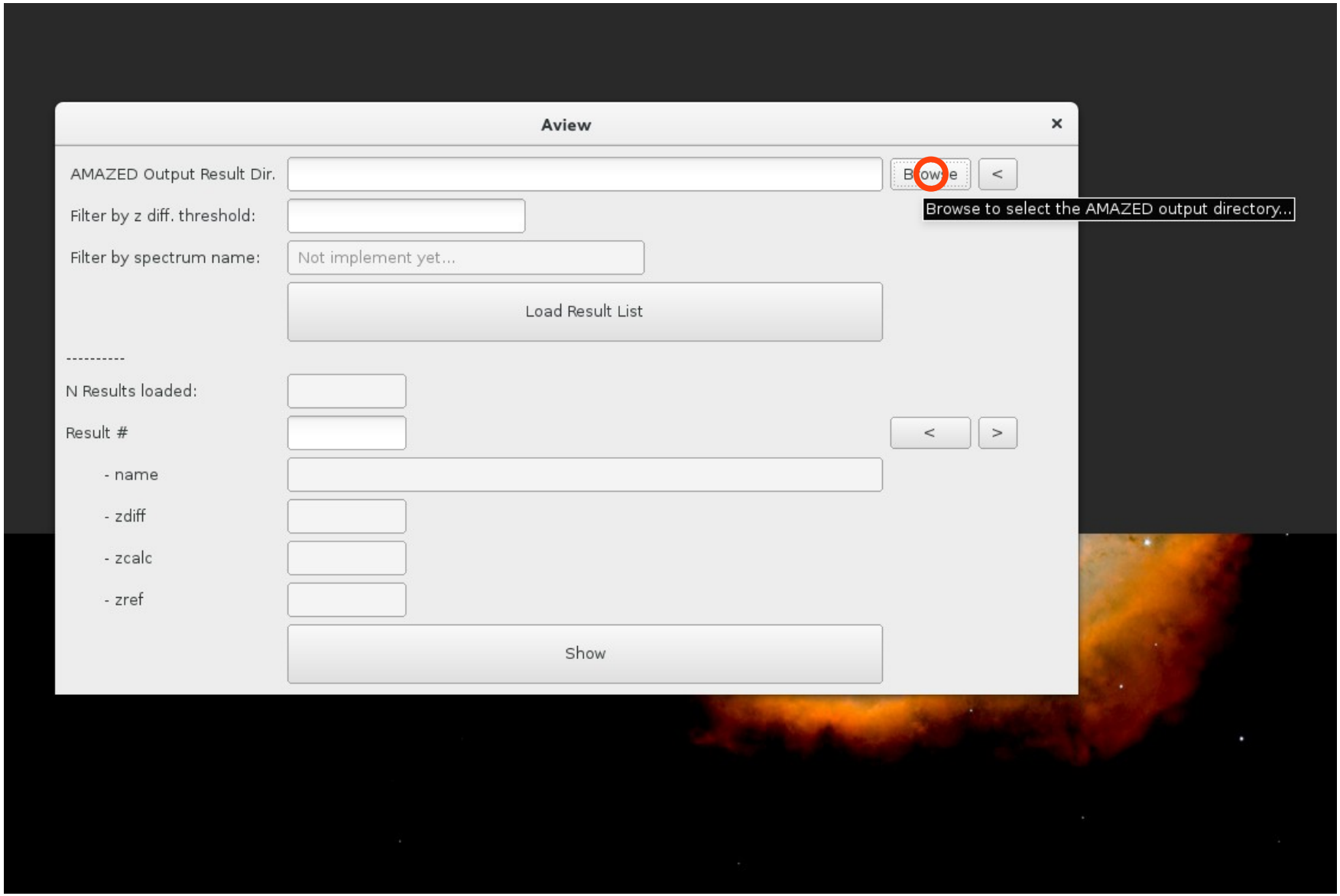
aviewgui.py

Quick Start
2016/01/05

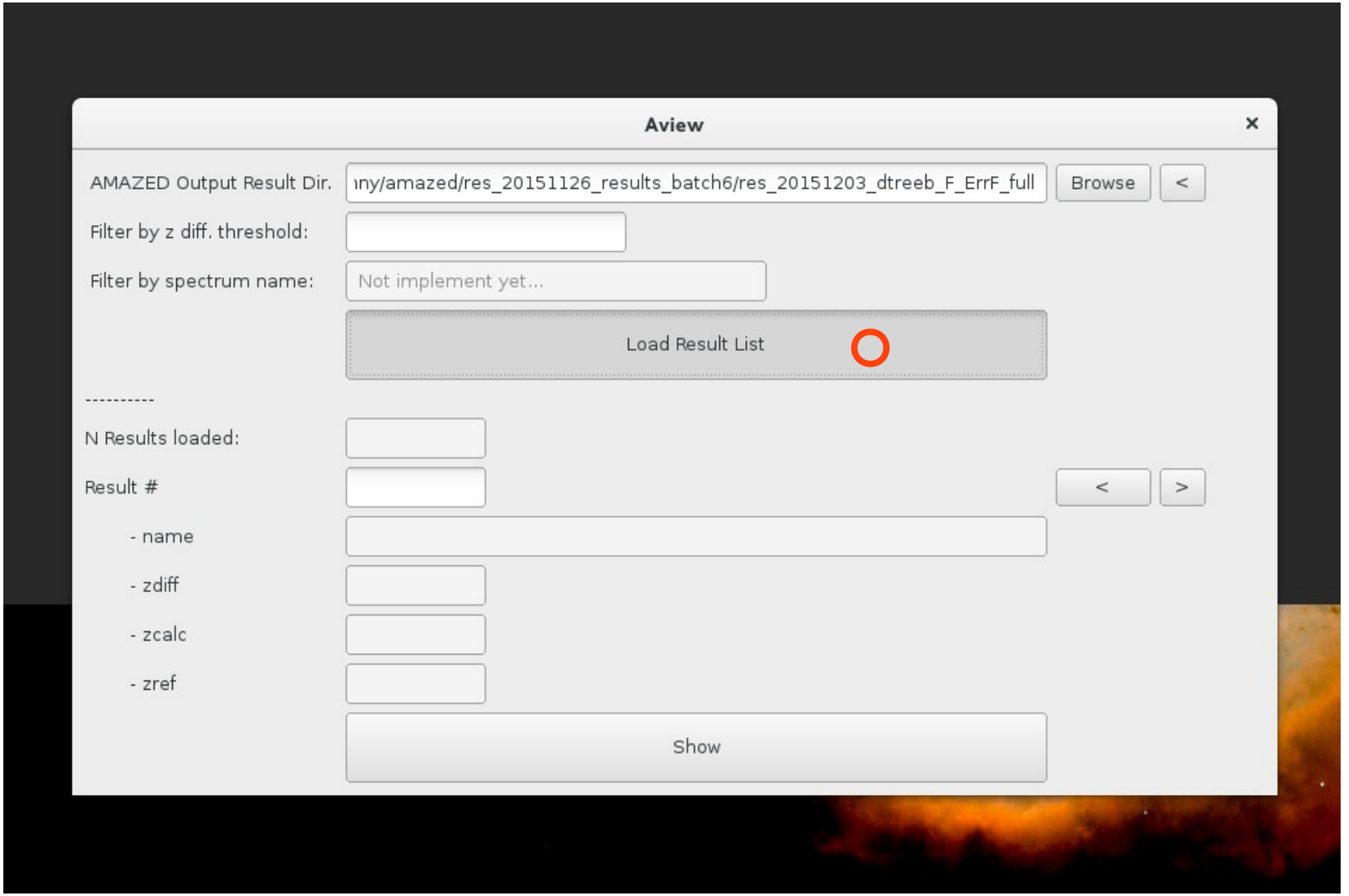
Start the script



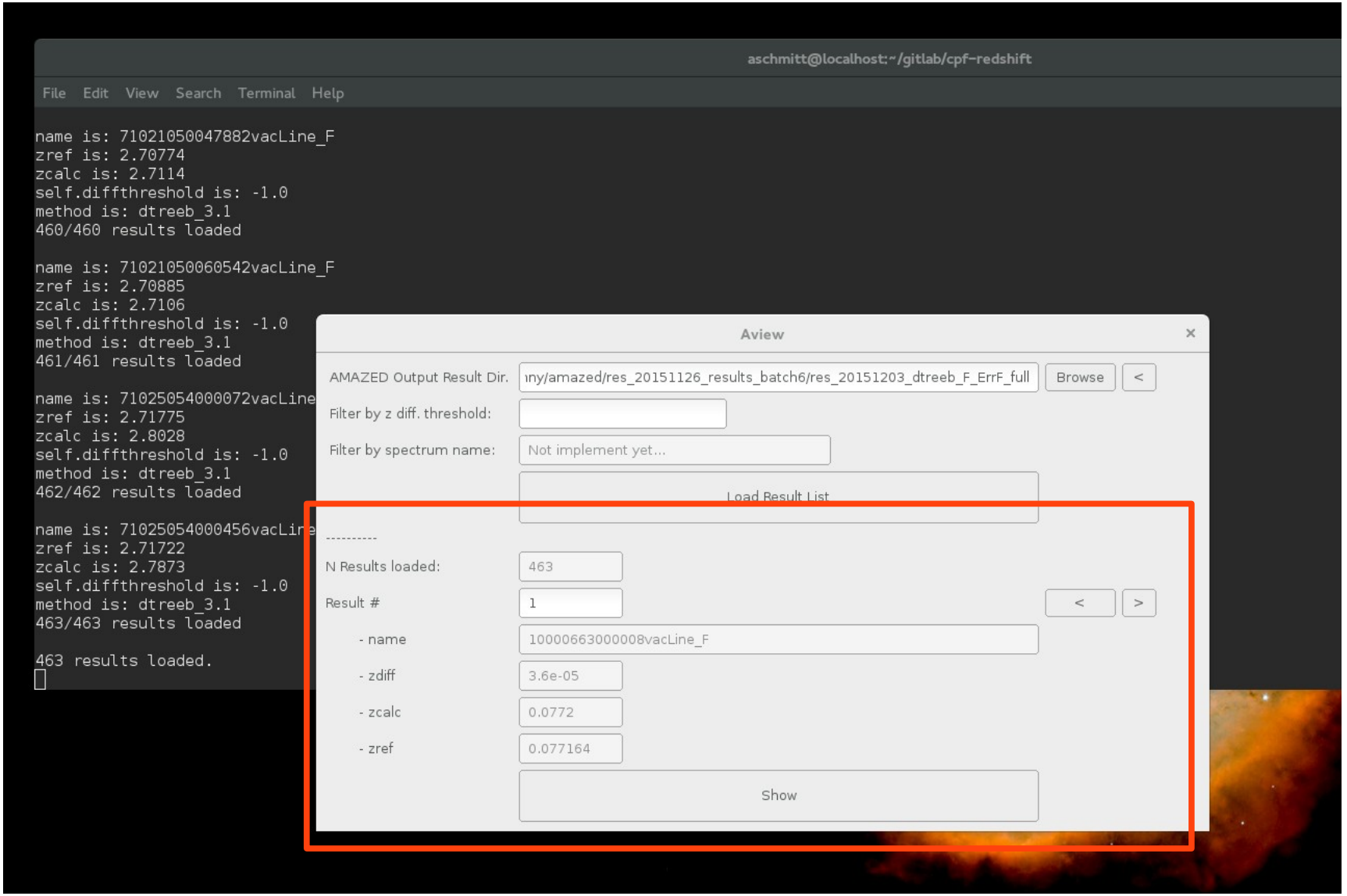
Browse to the AMAZED output directory



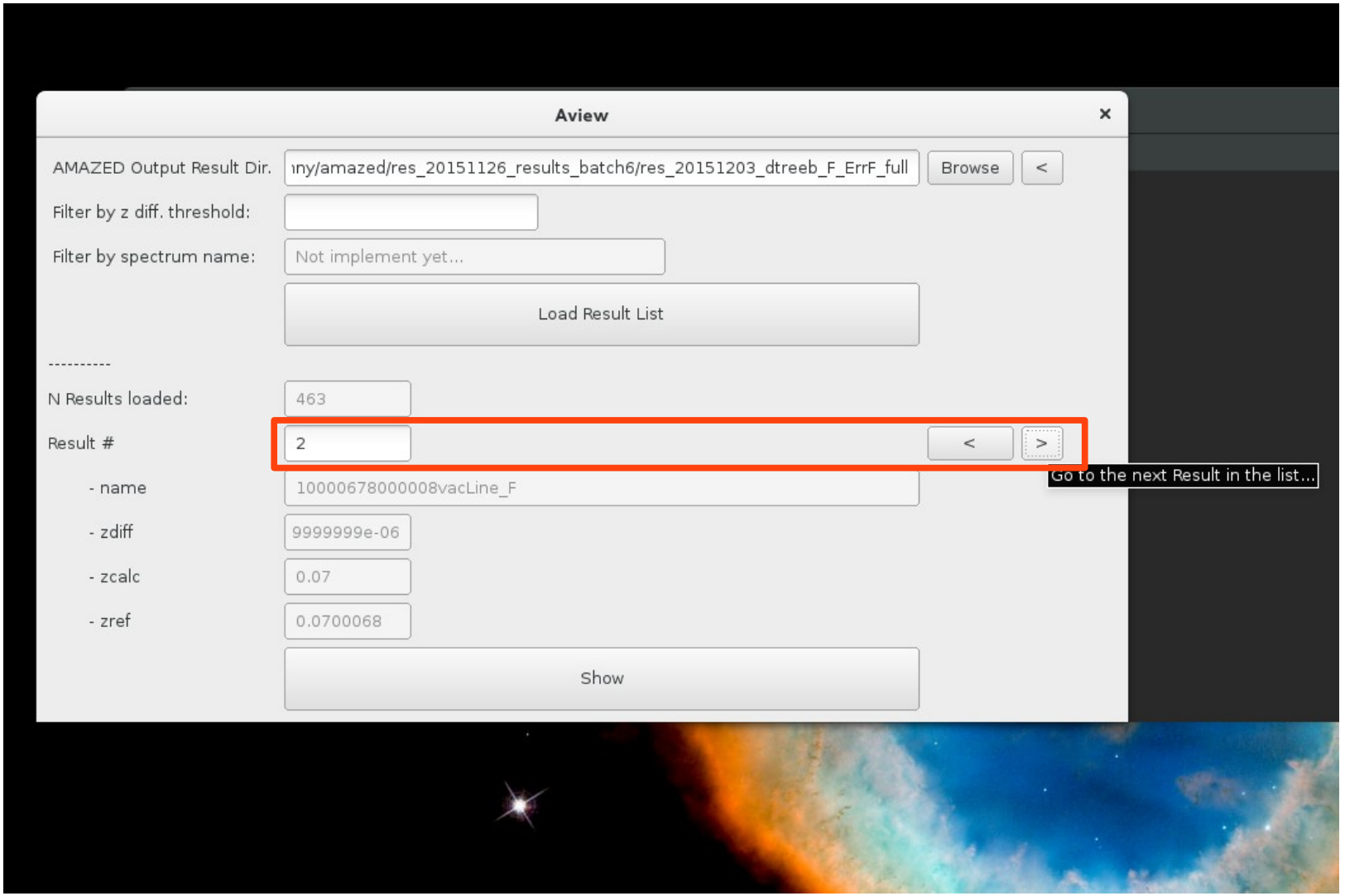
Load the result list



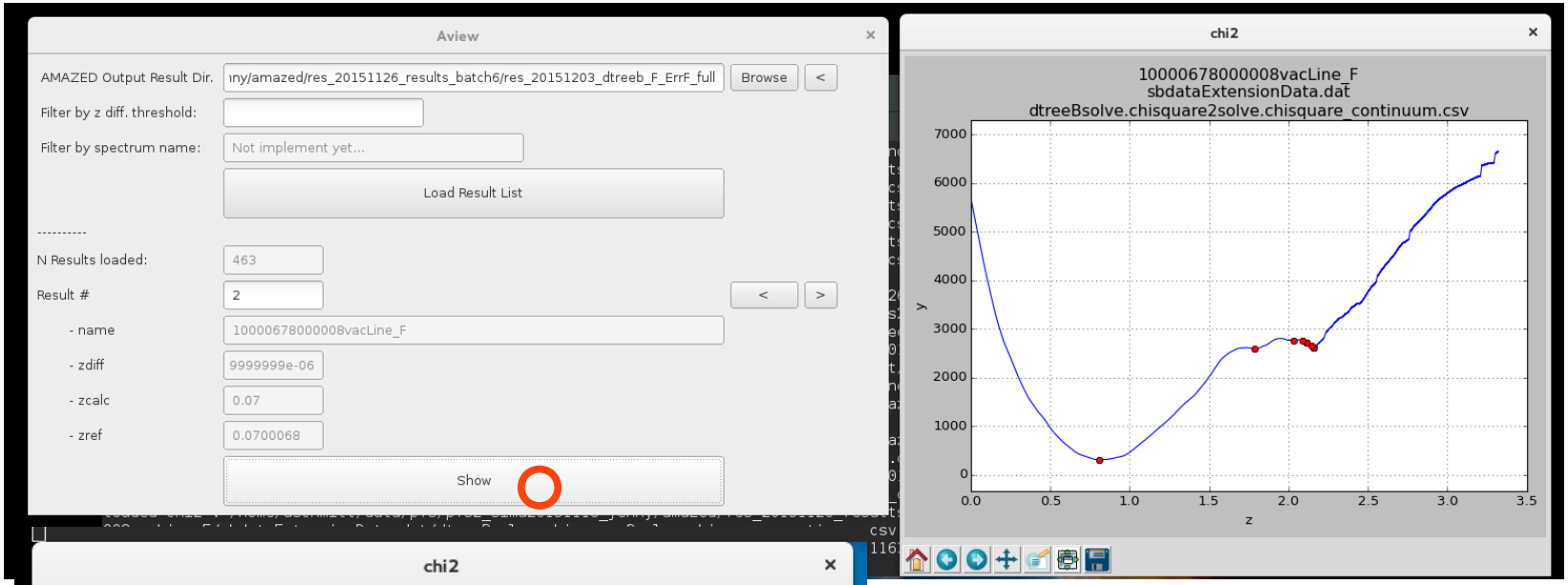
See the loaded results...

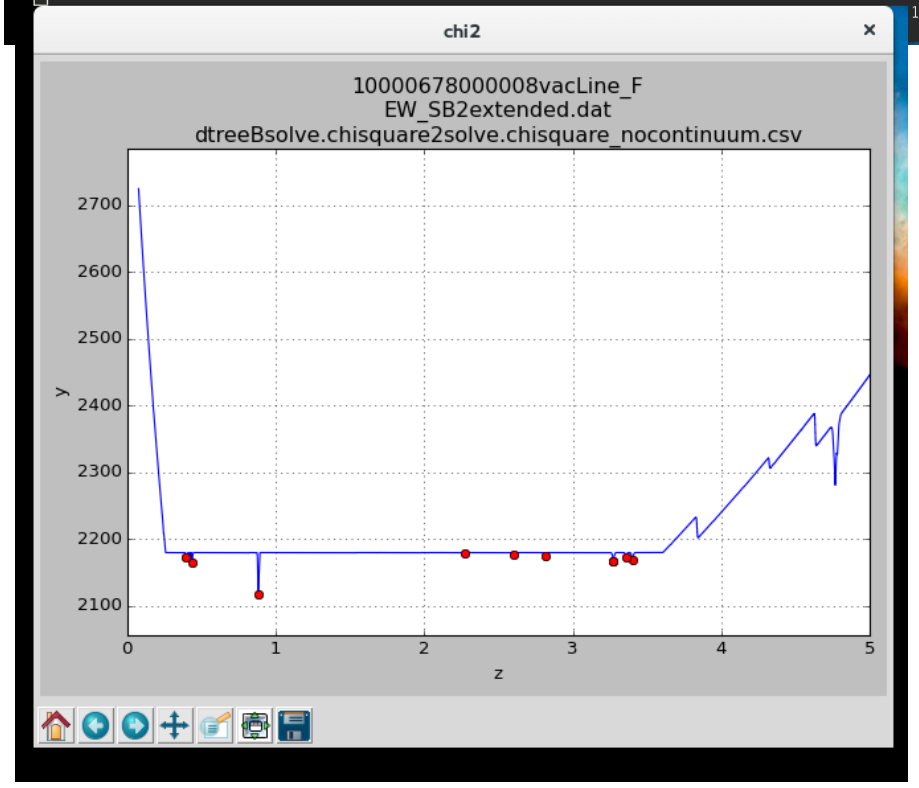


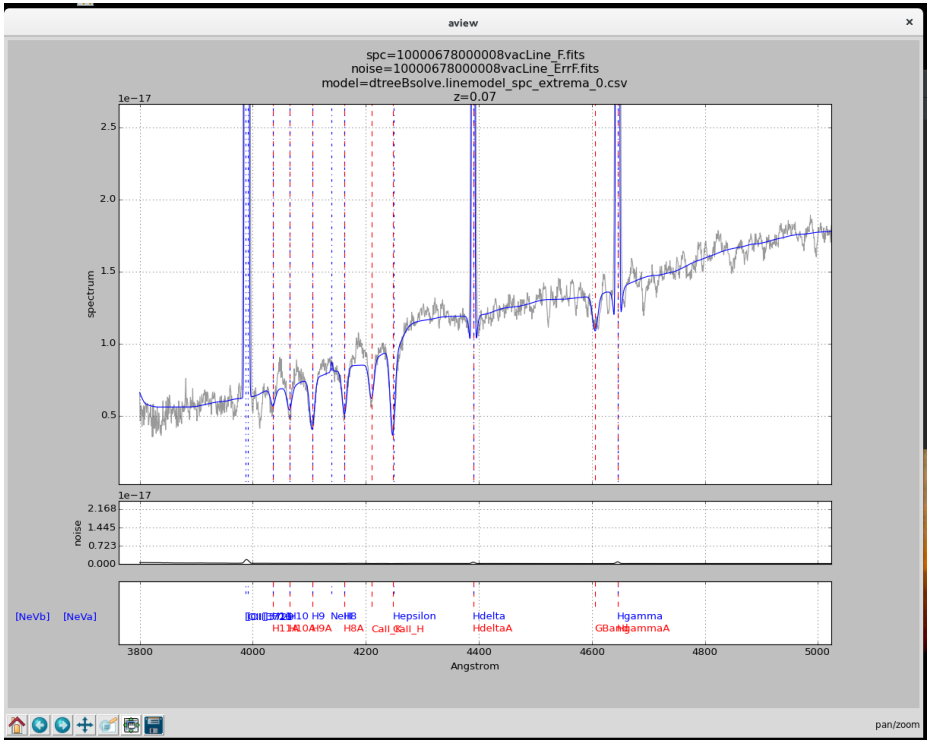
Navigate between the individual spectrum results...



Plot some results for the selected spectrum ...







NB:

- You can use aview.py for a command line version of the display script. It allows to override the redshift value (default is the *zcalc* value) used for plotting the spectrum overlaid with the template or the linemodel…
- The chisquare results are only displayed if the AMAZED pipeline has been run with the parameter :

  "SaveIntermediateResults": "all"
- Tested methods are : dtree7, dtreeB, chisquare, linemodel, ...
- This is an early prototype version that is intended to provide a quick and easy display. Your feedback is very welcome