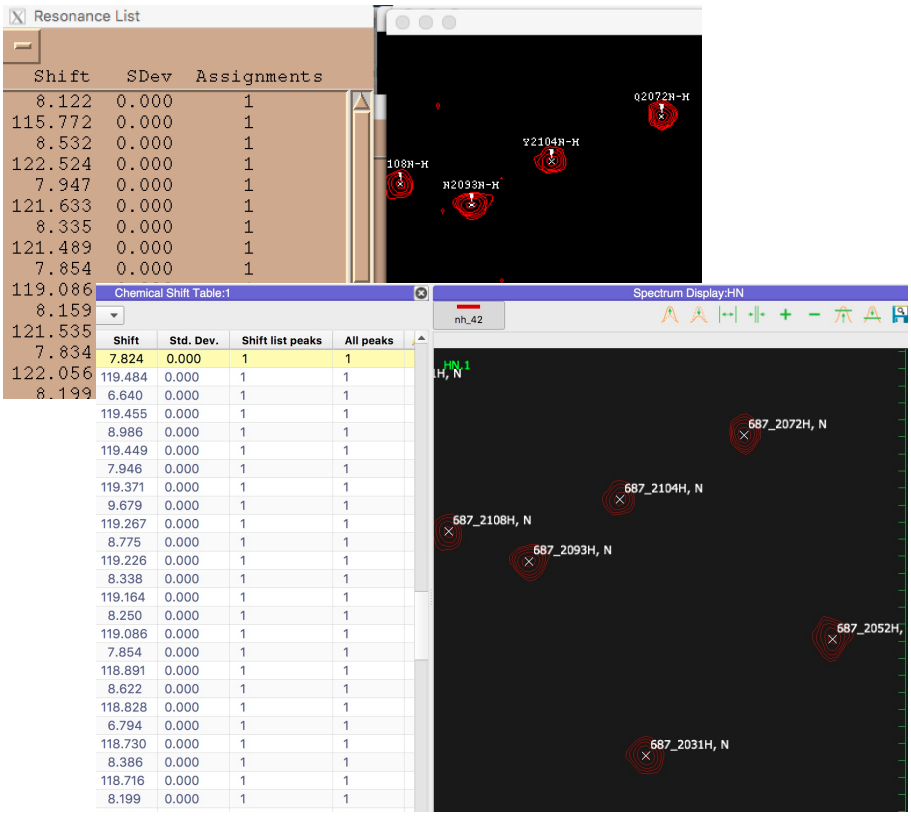
How To's:

Import Sparky project



Introduction

This How-To will show how to import a Sparky project directly to CcpNmr Analysis V3.

NB. In the current release of Analysis, the sparky reader is still under development, we cannot guarantee that all Sparky projects will be opened correctly.

Start CcpNmr Analysis V3

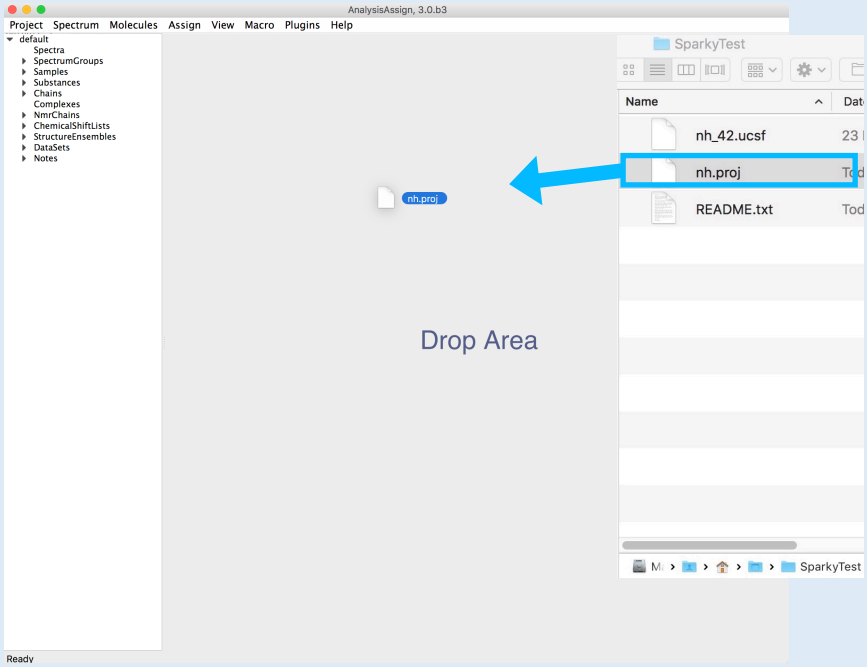
Apple users by double clicking the icon *CcpNmrAnalysis*

Linux users by using the terminal command: *bin/assign*

How To 1

Import Sparky

Drop nh.proj



1A **Drop a *.proj file**

- Find a sparky project on SparkyTest directory
- Simply Drag and drop the file **nh.proj** onto the sidebar or Drop Area

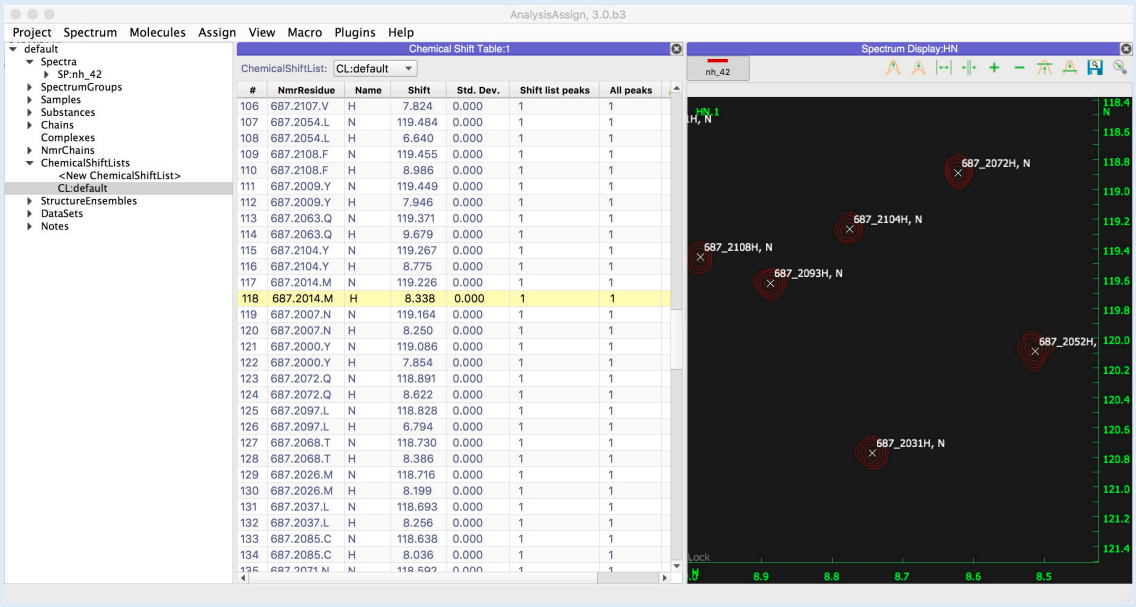
The project should be fully loaded.

All the spectra and other data will be located in sidebar.

- Go on sidebar, expand the Spectra branch
- Select the item SP:nh_42
- Drag and drop on the drop area

How To 1

Import Sparky



1B Open the Chemical Shift list

If you had an assignment in the sparky project, you may want to see the chemical shift list

- Go on sidebar, expand the ChemicalShiftLists branch
- Select the item CL:default
- Drag and drop on the drop area

Contact Us

Website:
www.ccpn.ac.uk

Suggestions and comments:
ccpnmr3@google.com

Issues and bug report:
https://bitbucket.org/ccpnmr/issue-tracker/

Cite Us

Skinner, S. P. et al. CcpNmr AnalysisAssign: a flexible platform for integrated NMR analysis. J. Biomol. NMR 66, (2016)

Tutorial Version History:

beta3 (LGM): First version