

# nMOLDYN

Computation and decomposition  
of **N**eutron scattering spectra  
from **M**OLEcular **D**YNamics  
simulations

# nMOLDYN 1 : a bit of history



ELSEVIER

Computer Physics Communications 91 (1995) 191–214

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Computer Physics  
Communications

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## *n*MOLDYN: A program package for a neutron scattering oriented analysis of Molecular Dynamics simulations

Gerald R. Kneller<sup>a,b,1</sup>, Volker Keiner<sup>c,2</sup>, Meinhard Kneller<sup>c,3</sup>, Matthias Schiller<sup>c,4</sup>

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Received 18 November 1994

# Original Fortran 77 version

# nMOLDYN 2 : a bit of history

## Software News and Updates

### ***n*Moldyn: A Program Package for a Neutron Scattering Oriented Analysis of Molecular Dynamics Simulations**

**T. RÓG,<sup>1,2</sup> K. MURZYN,<sup>1,2</sup> K. HINSEN,<sup>2</sup> G. R. KNELLER<sup>2</sup>**

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45071 Orléans Cedex 2, France*

*Received 30 July 2002; Accepted 1 November 2002*

Version 2:

Fortran → Python

Inclusion of MMTK library (Hinsen)

J Comp Chem 24: 657–667, 2003

# nMOLDYN 3 : a bit of history



Journal of  
**COMPUTATIONAL  
CHEMISTRY**

WWW.C-CHEM.ORG

SOFTWARE NEWS AND UPDATES

## *n*Moldyn 3: Using Task Farming for a Parallel Spectroscopy-Oriented Analysis of Molecular Dynamics Simulations

Konrad Hinsen,<sup>[a,b]</sup> Eric Pellegrini,<sup>[c]</sup> Sławomir Stachura,<sup>[a,b]</sup> and Gerald R. Kneller\*<sup>[a,b,d]</sup>

Version 3:

ILL involvement – Eric Pellegrini

Major code upgrade

Task-farming parallelisation (Pyro)

J of Comp Chem 2012, 33, 2043–2048

# nMOLDYN @ ILL (Eric Pellegrini)



## Version 3:

Agence National pour la Recherche (ANR) project –  
THALER (2008 – 2010 inc)

Post-doc: Eric Pellegrini

## Version 4 (MDANSE):

ANR project – SPUTNIK (2011 – 2013 inc)

Post-docs: Bachir Aoun (2 yrs), Gael Goret (1 yr)

Build server/CI: Jenkins with Yannick Raoul, GitLab  
with Jamie Hall & Fabien Pinet (ILL/IT)

Scientific support: with Miguel Gonzalez

All supervised by Eric Pellegrini



# nMOLDYN 3

## **THALER Objectives:**

- Improving portability → easier installation ...
- Reorganization of the code
- New scientific features
- New Graphical User Interface
- Improving readability → developer/user
- Improving ergonomomy
- Improving performance → parallelization

# nMOLDYN 3 – new scientific features

SCATTERING:

DYNAMICS:

STRUCTURE:

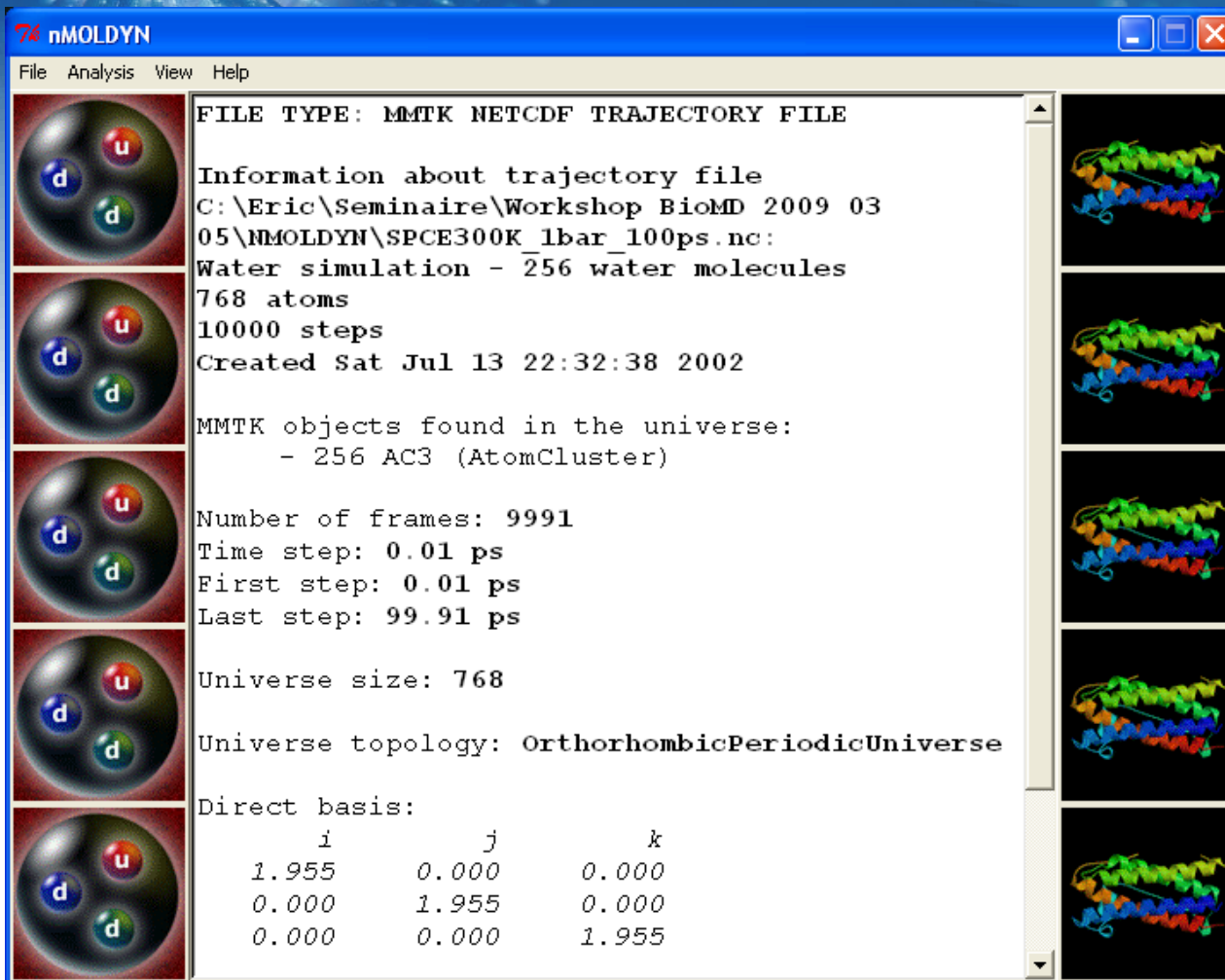
NMR:

TRAJECTORY:

- Center Of Mass Reduced Trajectory
- Global Motion Filtered Trajectory
- Quasi Harmonic Analysis
- ScrewFit Analysis (Paolo)
- Reduced Trajectory

Read trajectories from many MD codes (classic & DFT): CHARMM, NAMD, LAMMPS, DL\_POLY, (AMBER, GROMACS ?) VASP, Materials Studio (DISCOVER, FORCITE, CASTEP, DMOL, DFTB)

# nMOLDYN 3 - new GUI



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File Analysis View Help

FILE TYPE: MMTK NETCDF TRAJECTORY FILE

Information about trajectory file  
C:\Eric\Seminaire\Workshop BioMD 2009 03  
05\NMOLDYN\SPCE300K\_1bar\_100ps.nc:  
Water simulation - 256 water molecules  
768 atoms  
10000 steps  
Created Sat Jul 13 22:32:38 2002

MMTK objects found in the universe:  
- 256 AC3 (AtomCluster)

Number of frames: 9991  
Time step: 0.01 ps  
First step: 0.01 ps  
Last step: 99.91 ps

Universe size: 768

Universe topology: OrthorhombicPeriodicUniverse

Direct basis:

	<i>i</i>	<i>j</i>	<i>k</i>
	1.955	0.000	0.000
	0.000	1.955	0.000
	0.000	0.000	1.955



# nMOLDYN 3 - new GUI

**subset selection dialog**

Settings

Selection media

from a selection file     from the loaded trajectory     from an expression string

Selection from the loaded trajectory

Object name	Selection keyword	Selection value
P892	chemfragment	hydroxy

Object name: P892

Selection keyword: chemfragment

Selection value: hydroxy

Linkers

Clear Undo ( ) AND OR

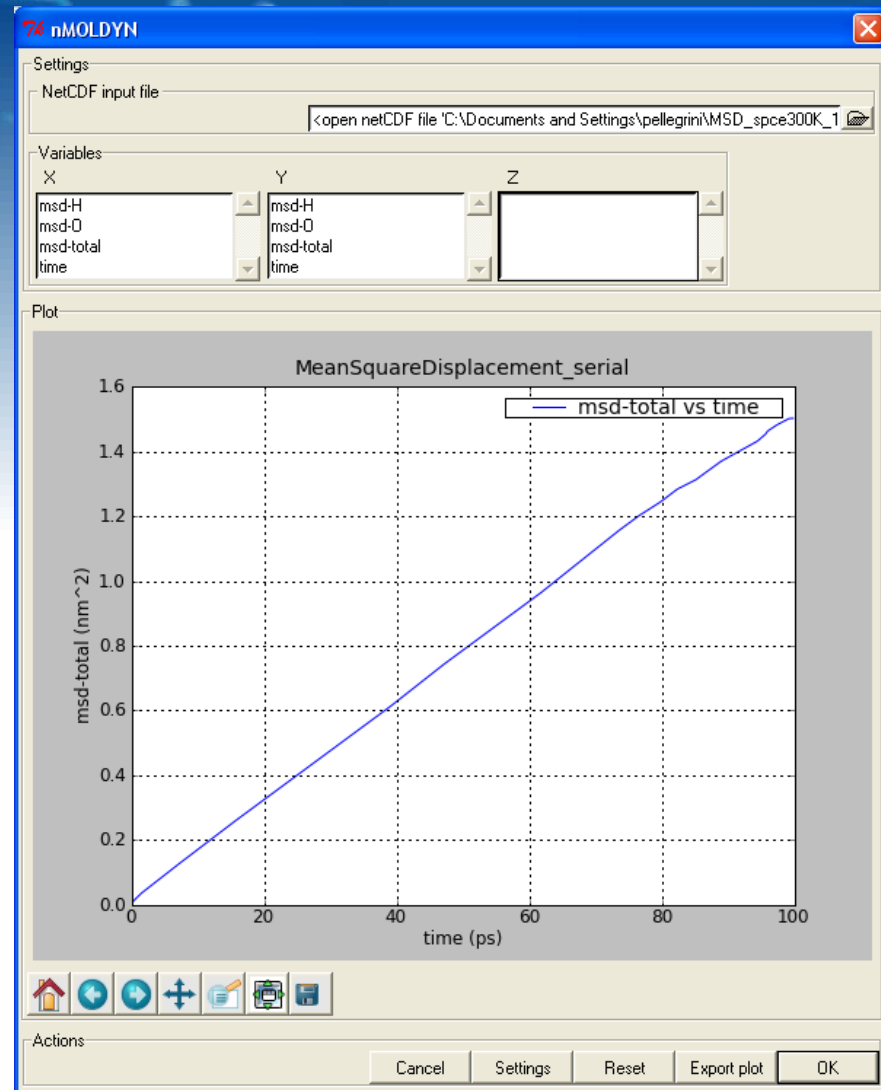
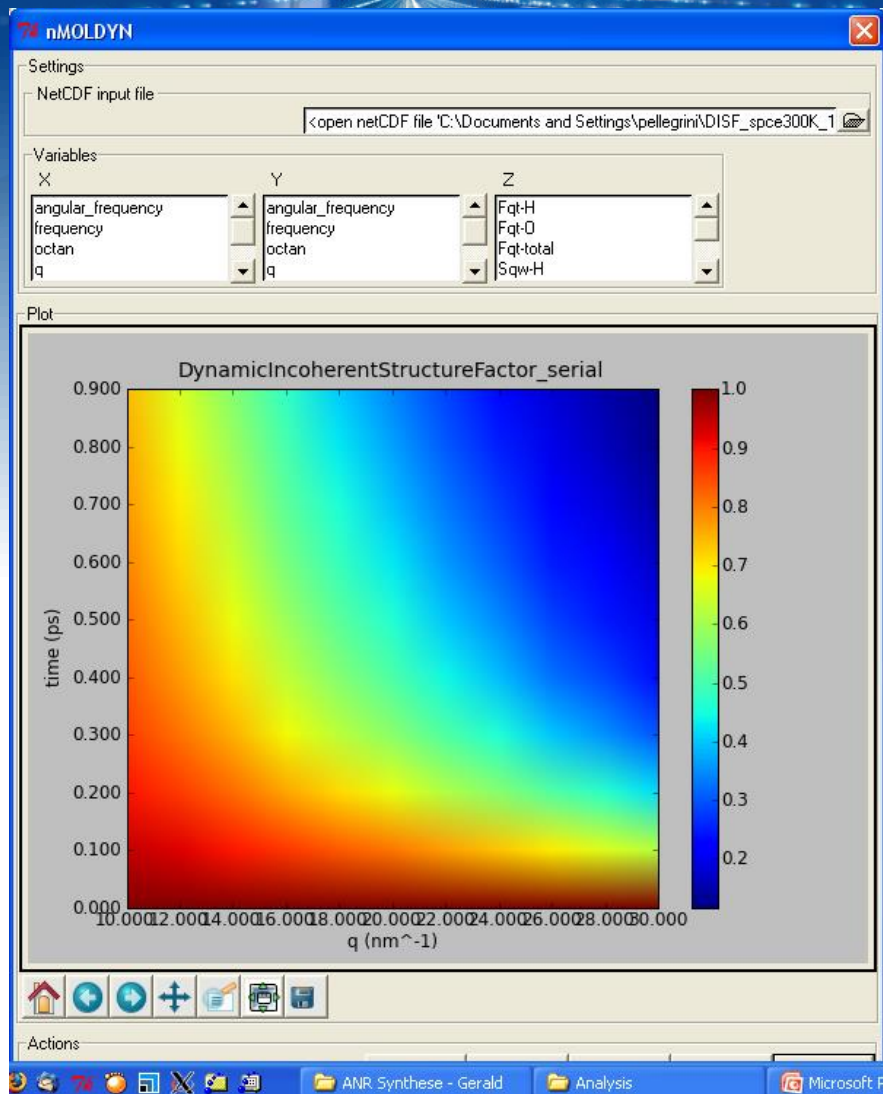
Selection preview

```
object objectname P892 chemfragment hydroxy AND chemfragment methyl
```

Actions

Cancel OK

# nMOLDYN 3 - new GUI



# nMOLDYN 3 - new GUI

**Mean-Square Displacement** [X]

Setup

Trajectory file  
C:/Eric/nMOLDYN/Trajectories/NetCDF/spce300K\_1bar\_100ps\_paolo.nc

Frame selection (14.101 ns to 114.001 ns step 0.1 ns)  
1:9991:10

Project displacement on  
no

Subset selection  
all [Select]


Deuteration selection  
no [Select]

Weights  
 equal     mass     coherent  
 incoherent     atomicNumber

Pyro server  
monoprocessor [Select]

Output file  
C:\Documents and Settings\pellegrini\MSD\_spce300K\_1bar\_100ps\_paolo.r [Browse]

Actions  
[Cancel] [Save] [Run] [Save/Run] [Estimate]

Progress (in %)  
100  


## **SPUTNIK Objectives:**

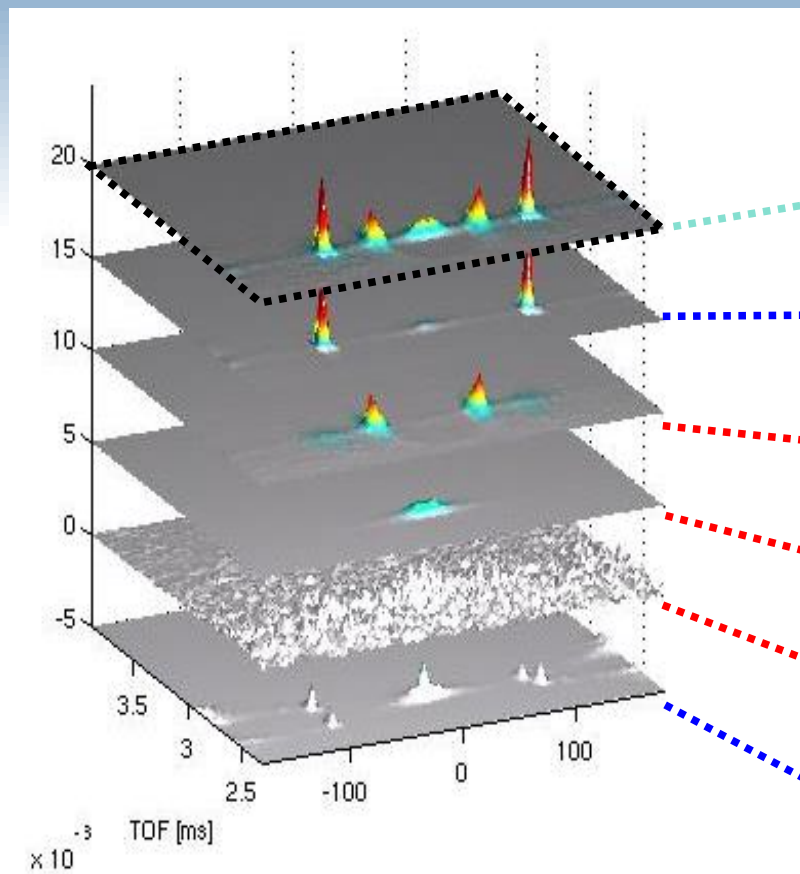
- Integrating experiment related effects – neutron flight simulator
- « Combining » nMoldyn and McStas
  - Simple interface between the codes
  - Web service
- Integrating simple ray-tracing simulations in nMoldyn
- and/or analytical solutions
- On-going upgrade of nMoldyn (e.g. Linux/Mac install, 64-bit code, parallelisation/cluster optimisation, complete separation of front (GUI) and back end,...)

# Instrument simulations and virtual experiments





# Instrument simulations and virtual experiments



Total signal

Container (Nb)

Coherent (I-Rb)

Incoherent (I-Rb)

Multi. Scatt. \*100

Cryo-furnace (Al) \*10

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nMoldyn

File View Help

Plugins

- Miscellaneous
- K vectors
- Viewer
  - Molecular Viewer
- Analysis
  - Trajectory
  - Thermodynamics
  - Virtual Instruments
  - Dynamics
  - Scattering
    - Dynamic Coherent Structure
    - Elastic Incoherent Structure
    - Current Correlation Function
    - Gaussian Dynamic Incoherent Structure
    - Dynamic Incoherent Structure
- Structure
  - Static Structure Factor
  - Eccentricity
  - Coordination Number
  - Root Mean Square Deviation
  - Voronoi
  - Root Mean Square Fluctuation
  - Molecular Trace
  - Solvent Accessible Surface
  - Pair Distribution Function
  - Radius of Gyration
  - Density Profile
  - Spatial Distribution

3

single\_wall\_nanotube.nc protein\_in\_periodic\_universe.nc

Molecular Viewer

1

Data

- mmtk\_trajectory
- single\_wall\_nanotube.nc
- waterbox\_in\_periodic\_universe.nc
- protein\_in\_periodic\_universe.nc

2

Animation

30 95

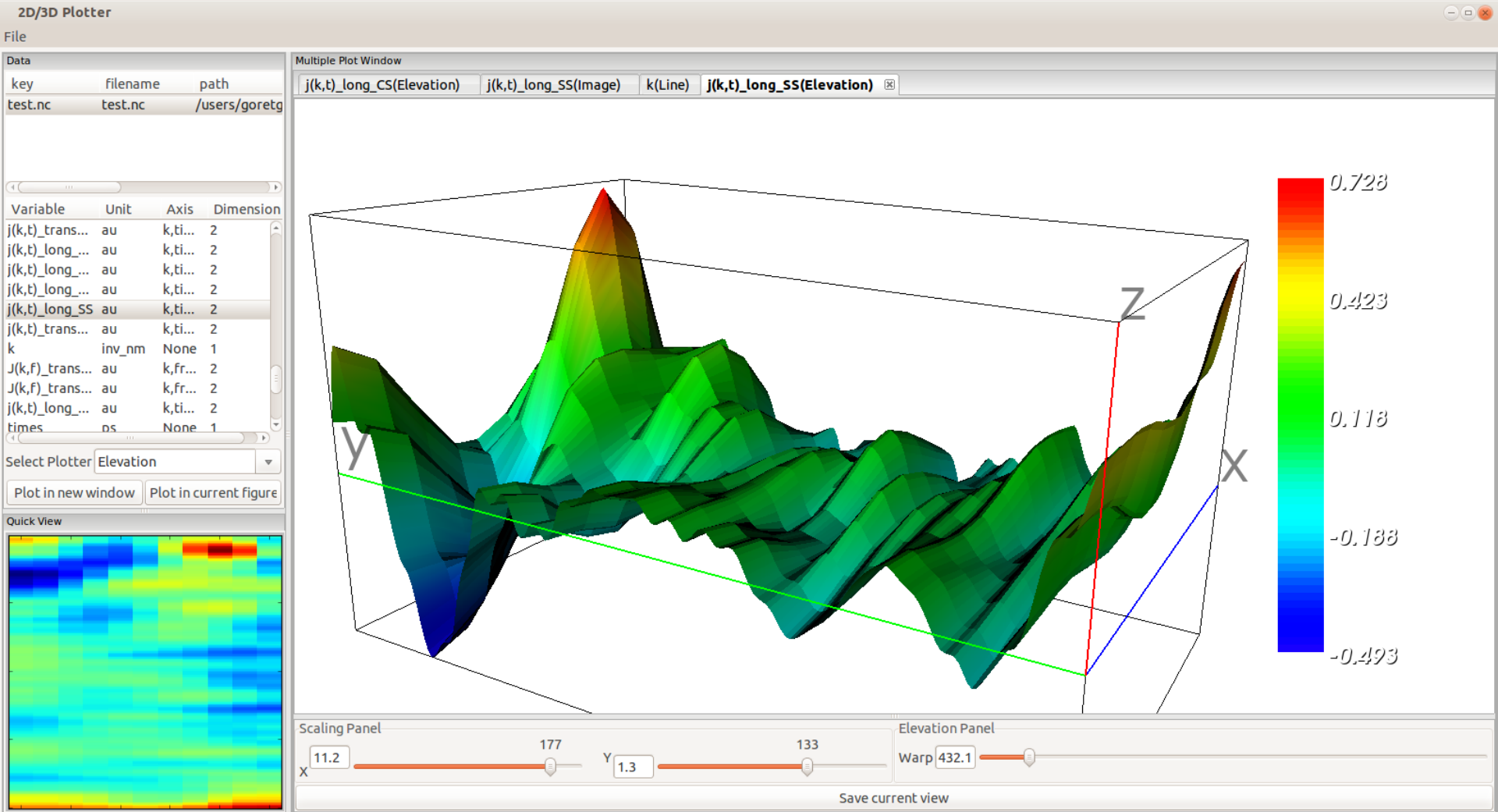
Logger Console Jobs

NAME	PID	START	ELAPSED	STATE	PROGRESS	ETA	KILL
5223_3wkz	5223	04-06-2014 09:57:32	00:00h:00m:16s	finished		N/A	
5249_drds	5249	04-06-2014 09:59:03	00:00h:00m:16s	finished		N/A	
5412_1qd2	5412	04-06-2014 10:09:15	00d:00h:00m:54s	running		00d:00h:00m:06s	

4



# nMOLDYN 4



4

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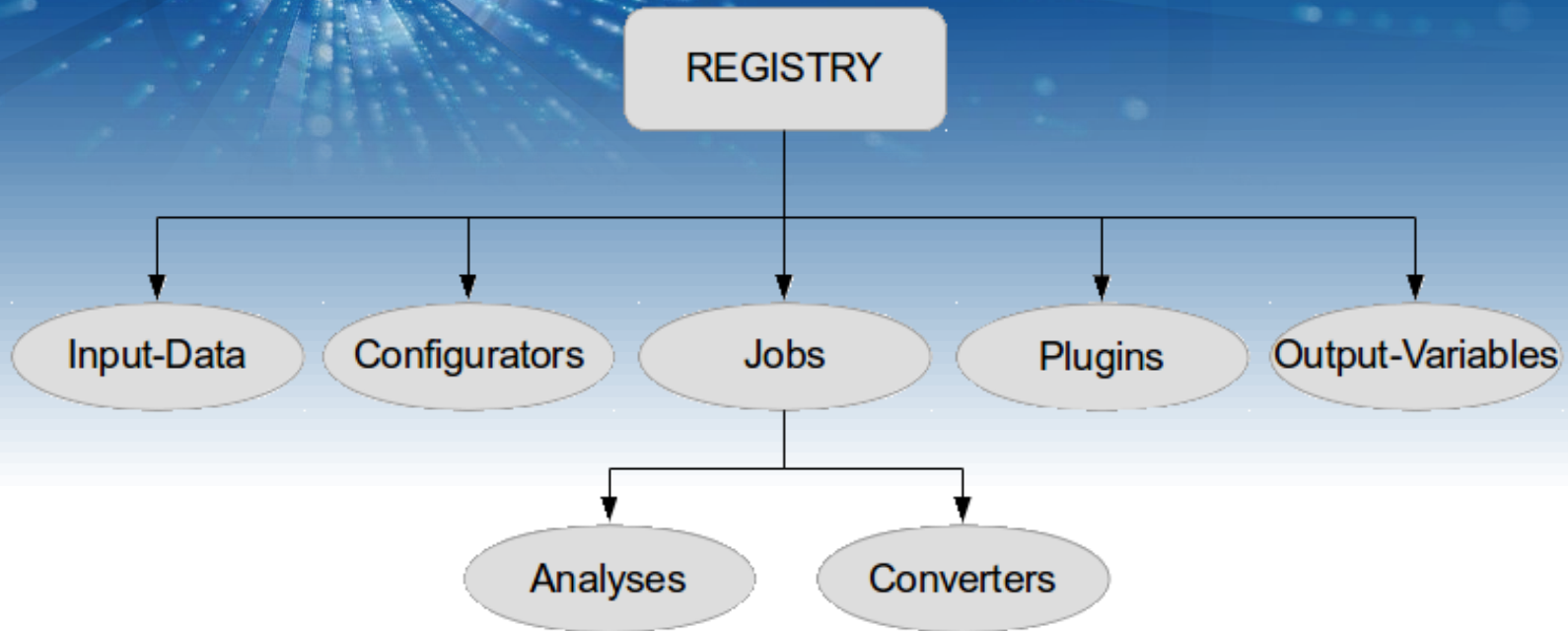
The screenshot displays the nMoldyn software interface. The main window shows a 3D visualization of neutron scattering paths, with a central source point emitting rays that pass through a sample and are detected by a detector array. The paths are color-coded in pink and cyan. A 'McStas Virtual Instrument Viewer' window is open, showing simulation parameters:

- mcstas options
  - neutron count: 10000
  - McStas output directory: /nm\_wdir/mcstas\_output/24.03.2014-
- trace the 3D view of the simulation:  Yes
- instrument parameters
  - beam wavelength Angs: 2.00000
  - environment thickness m: 0.00200
  - beam resolution meV: 0.10000
  - sample height m: 0.03000

The bottom of the interface features a 'Jobs' table with the following data:

Name	Pid	Owner	Start	State	Progress	Kill
gore_17397_25gsyp	17397	goretg	Mon Mar 24 15:37:10 2014	finished	<div style="width: 100%; height: 10px; background-color: blue;"></div>	<input type="button" value="X"/>

# nMOLDYN 4 – technical



Based on Python, Cython, WxWidget, Matplotlib, VTK,...

Automated GUI generation and menu update for new analyses



# nMOLDYN 4 → MDANSE

MDANSE – Molecular Dynamics to Analyse Neutron  
Scattering Experiments

nMoldyn

MMTK

# MDANSE 4 in SINE2020 ?

## SINE2020 Objectives: Idea...

- 1 person year of effort to include Lattice Dynamics in MDANSE
- Read vibration frequencies and displacement vectors (as a function of wavevector) from as many codes as possible
  - mainly DFT: VASP, CASTEP, DMOL, ABINIT, QUANTUM EXPRESSO, etc
  - or output from PHONOPY
  - alternative to PHONON from Krzysztof Parlinski, coupled to VASP, SIESTA, WIEN2K, widely used at ILL
- Analyse Lattice Dynamics
  - Calculate NS spectra for powders & single crystals on TAS (including TOSCA/Lagrange-like) and TOF
  - Provide additional analyses: thermodynamic quantities (enthalpy, entropy, free energy, heat capacity, MSD), participation ratio, Gruneisen parameters,...

## Nominally

- 2 person years of effort for MANTID @ ILL in addition to ILL Bastille/Endurance funding
  - ILL has recruited 3 programmers for 3 years starting May 2 2016
  - ILL has negotiated 1 year of consulting with Tessella (Ian Bush) starting April 18 2016)
- 1 person year of effort for QENS
- 1 person year of effort for Lattice Dynamics
- Not ready to start QENS and Lattice Dynamics because of MANTID activity
- ISIS/STFC (e-Science) starting a similar Lattice Dynamics project –combine resources, decide on most appropriate, collaborative solution