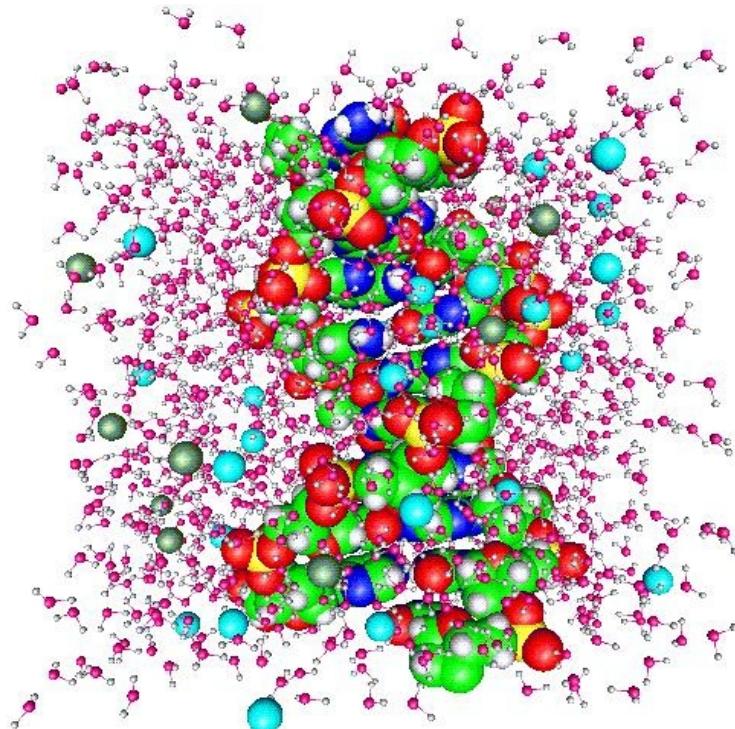


# Visualization of molecular simulations

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Molecular dynamics: solution of Newtonian equations of motion in the molecular system

For each atom at each time:

$$v_i(t+1/2\Delta t) = v_i(t-1/2\Delta t) + F_i(\{x_j(t)\})/m_i \Delta t$$

$$x_i(t+\Delta t) = x_i(t) + v_i(t+1/2\Delta t) \Delta t$$

1998: MdynaMix - parallel scalable molecular dynamics for arbitrary molecular mixtures

[http://www.fos.su.se/physical/sasha/md\\_prog.html](http://www.fos.su.se/physical/sasha/md_prog.html)

# Current project: Interactive Visualisation

includes:

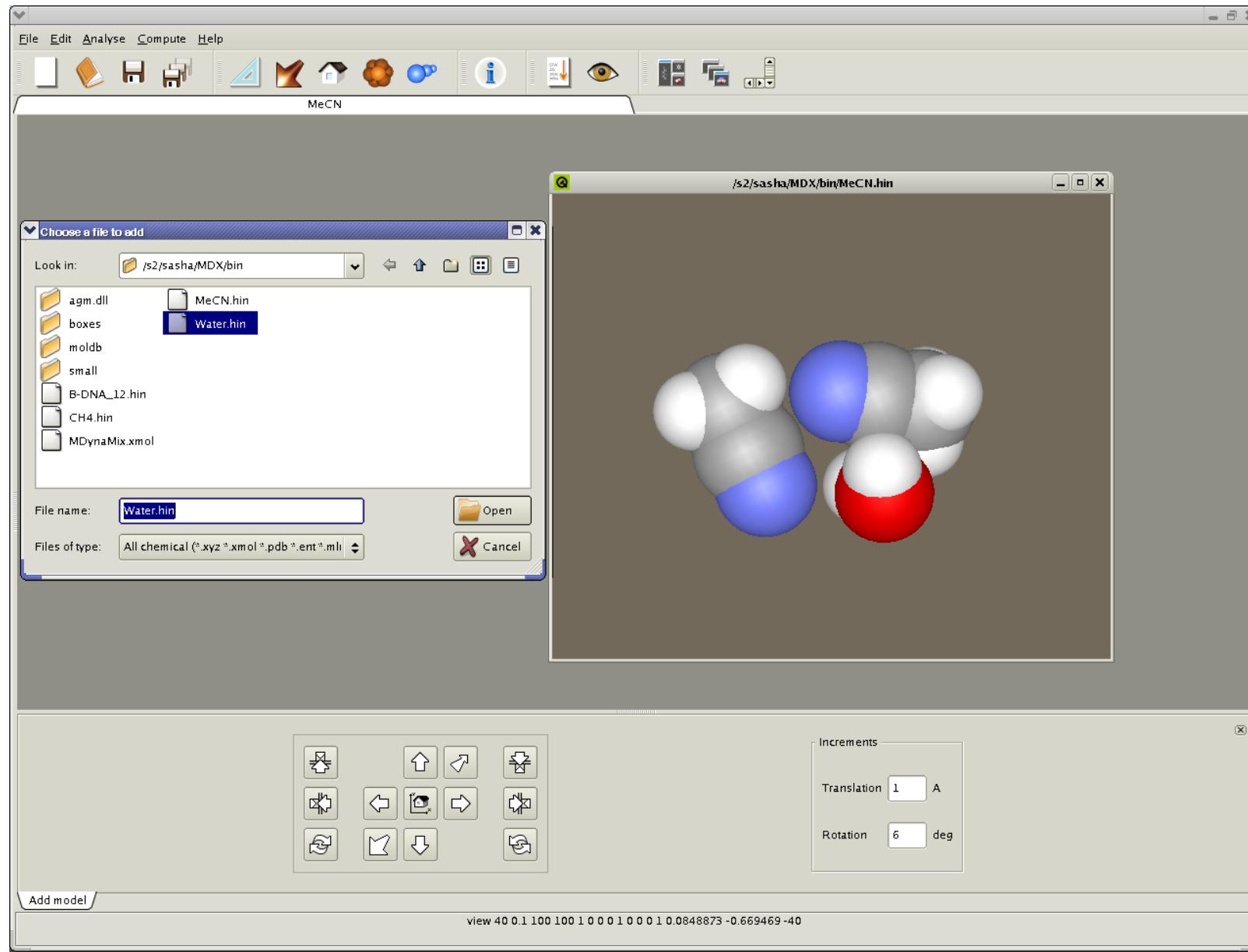
- initial preparing of molecular structures
- parameter (force field) association
- real-time visualization
- real-time analysis of result
- interactive: correction of simulation during the run

All these features are implemented in MDX project

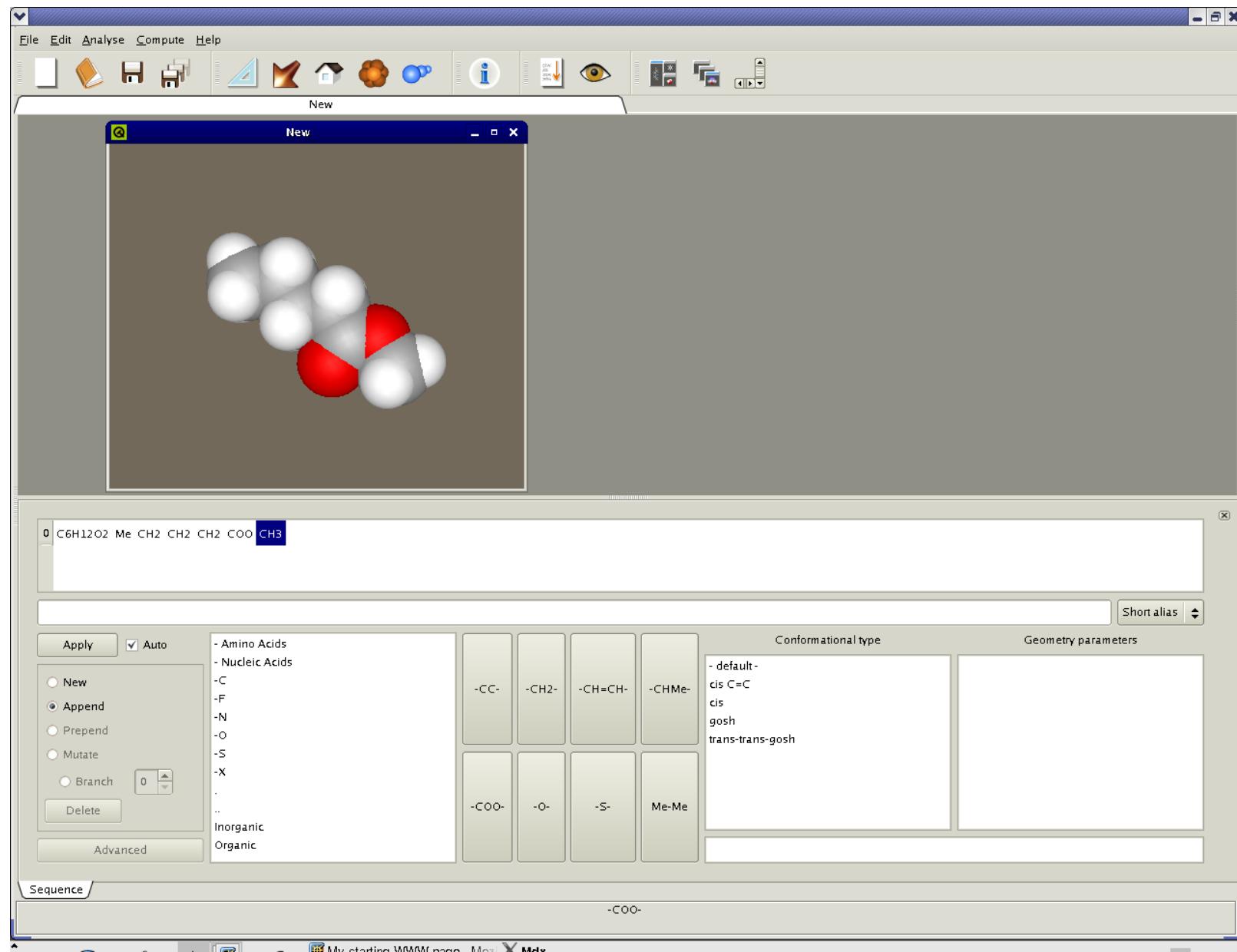
OS: Windows & Linux

# Snapshots from the program:

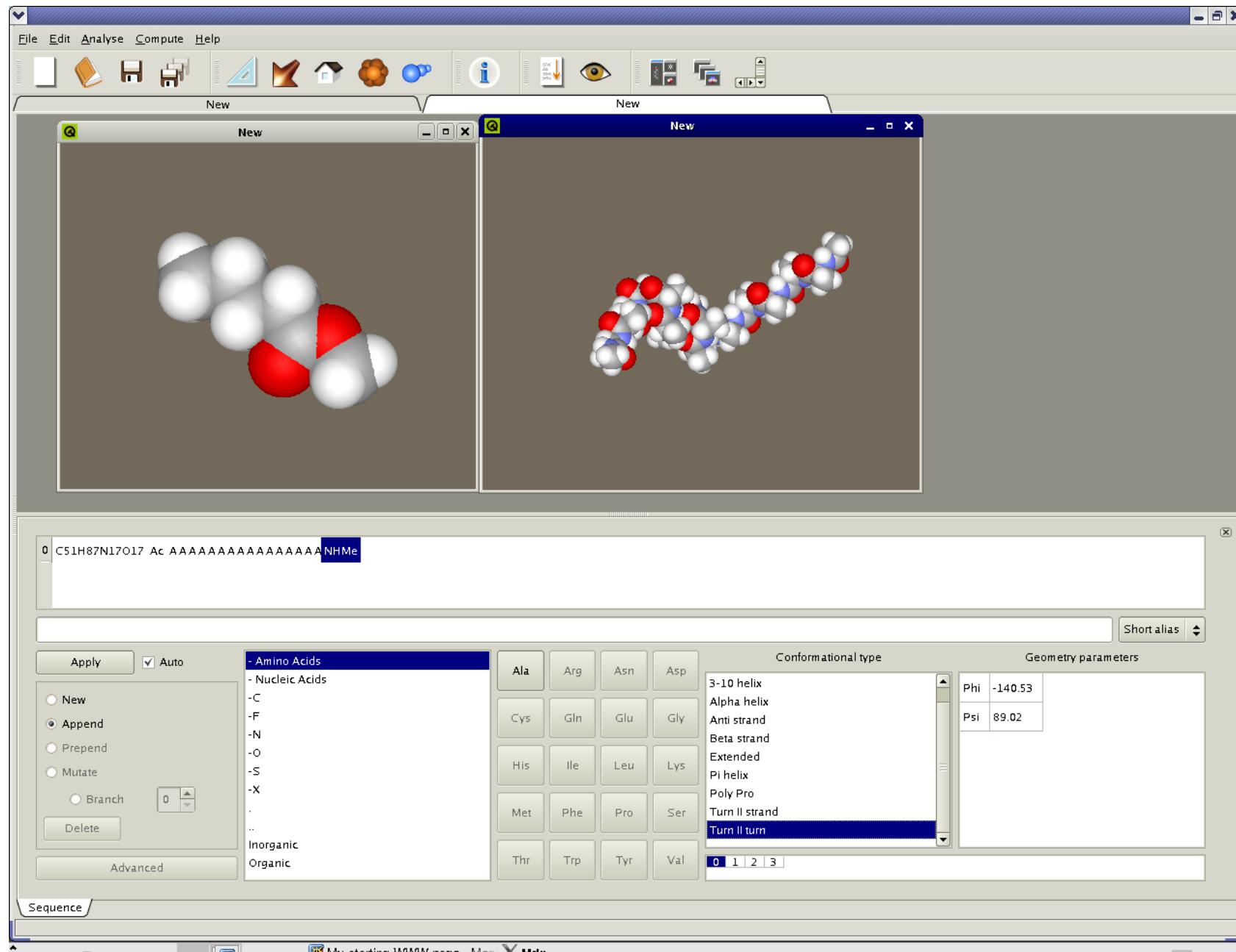
## Building system from available structures



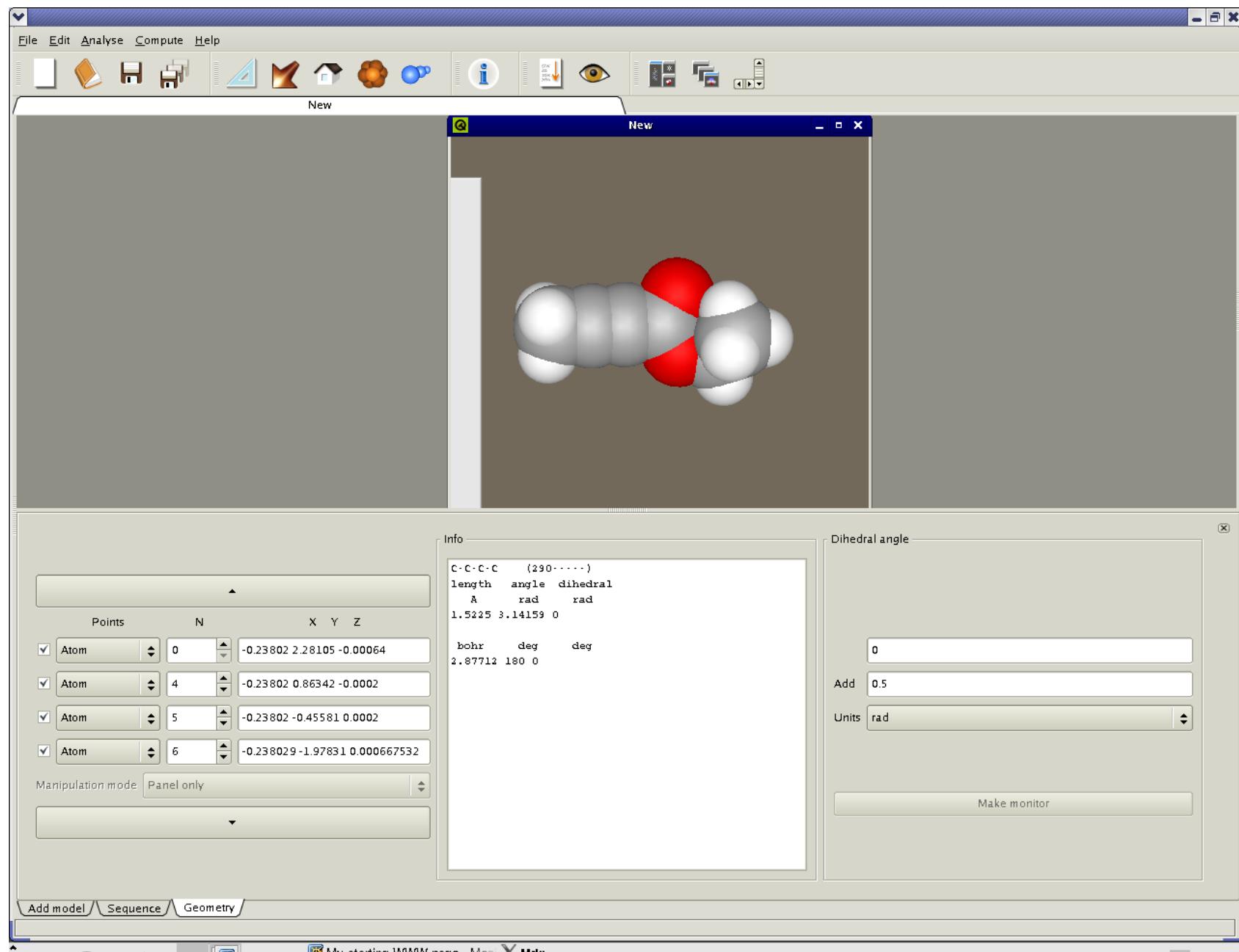
# Building arbitrary molecular structures



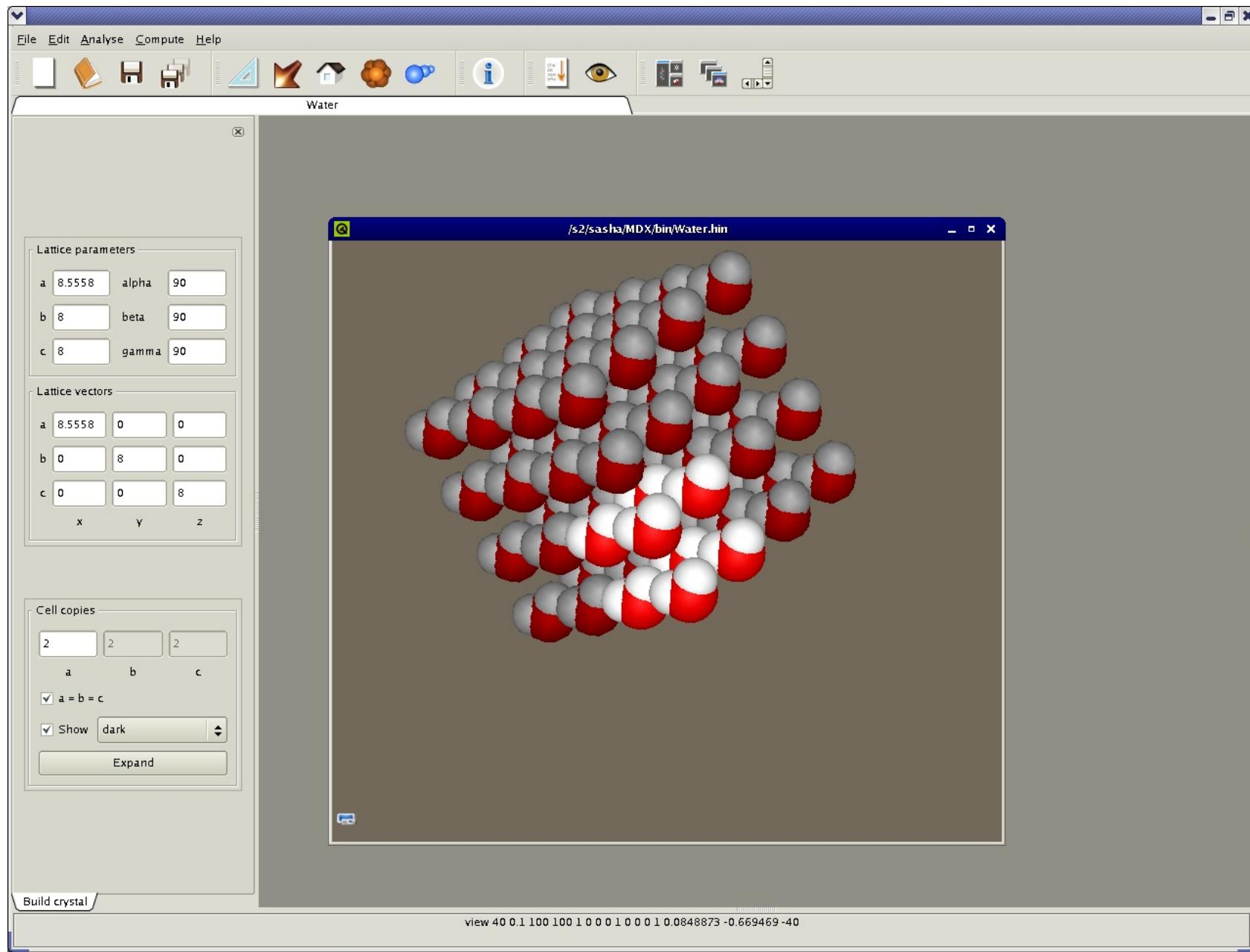
# Building proteins



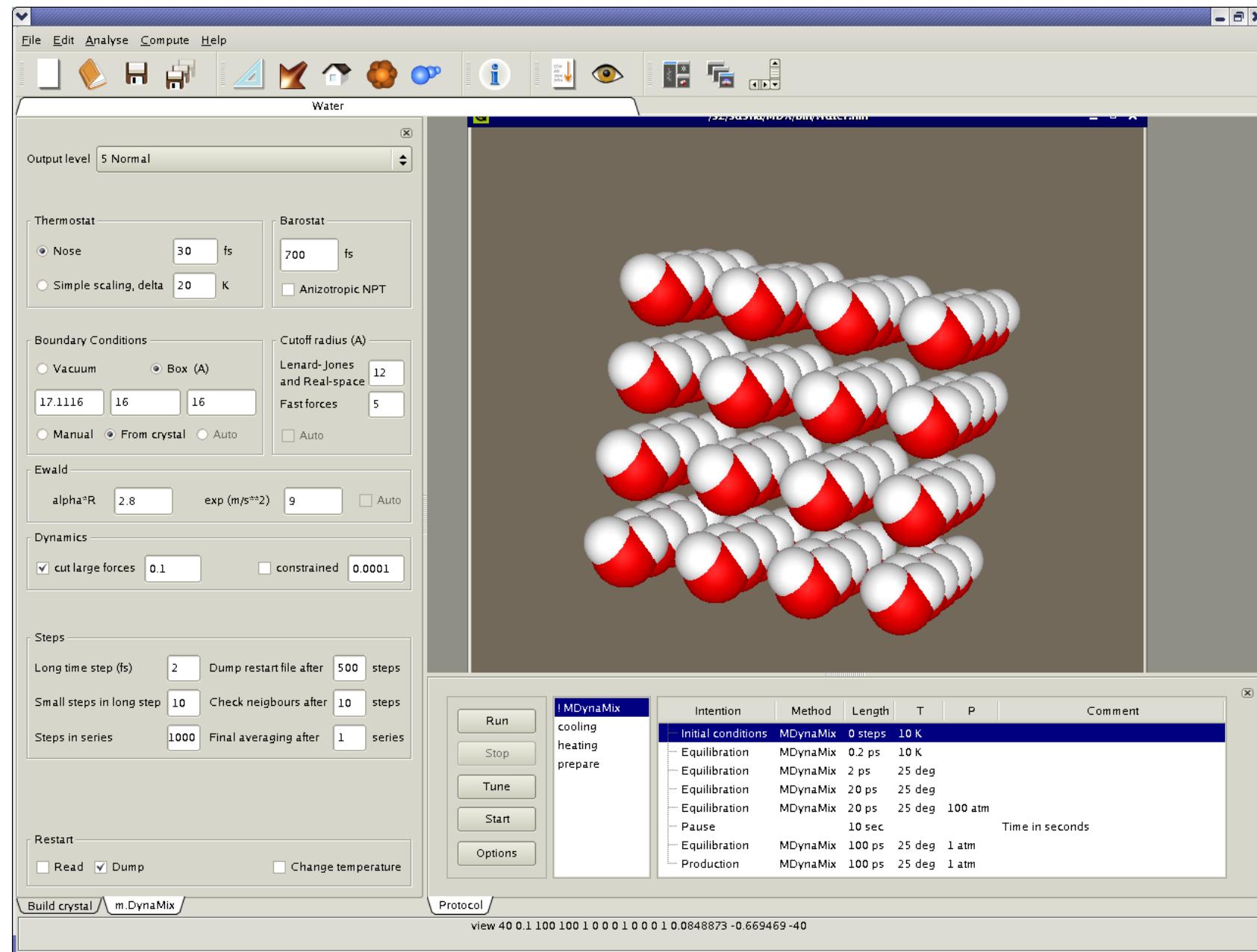
# Geometry editing



# Building a periodic system



# Control of molecular dynamics run



# Monitoring physical quantities

