

Ctrl+N Ctrl+O Ctrl+S H X F + - * Main toolbar

Selected Molecules

- All Ligands Ctrl-A
- All References Ctrl-R
- Protein Ctrl-P
- Model
- All Molecules Ctrl-D
- Selected Molecules Ctrl-M
- Picked Molecules

Change Display for...

- Show Fields as Spheres 0
- Show Fields as Tetrahedra 1
- Show Fields as Cubes 2
- Show Fields as Octahedra 3
- Show Fields as Dodecahedra 4
- Show Fields as Icosahedra 5
- Lines L
- Thin Stick T
- Capped Stick C
- Ball and Stick B
- CPK K
- Show/Hide Individual Fields...
 - Show Negative Field Points 6
 - Show Positive Field Points 7
 - Show Surface Field Points 8
 - Show Hydrophobic Field Points 9
- Field Shapes...
- Show Field Constraints
- Labels...
- Surfaces...
- Molecule Color
- Reset Display Ctrl+Shift+R
- Reset Display Completely

Right click

- Full Screen F11
- Select
- Copy This Molecule
- Export Selected
- Edit
- Edit a Copy
- Rename
- Delete This Molecule
- Delete This Alignment
- Set as Preferred Alignment
- Re-align This Molecule
- View Log
- Copy as New Reference
- Center on Picked Atom
- Reset View Center
- Save Image As...
- Capture Scene F4

Vertical toolbar shortcuts: P, A, R, I, Ctrl+G

Bottom status bar: Cpd342 0.945 Score 0.945 | Act 9.540 | 06 Activity Atlas for Potency[nM] Adenosine A3 receptor [Novelty Very High] | 94 molecules, 1 selected & 1 favs. Cpd342

Left mouse rotate, right mouse translate, middle mouse/wheel/Alt-left scales. Shift-mouse z-clip.

Shortcut	File operation
F1	Show manual
Ctrl+Shift+S	Save project as
Ctrl+S	Save project
Ctrl+L	View log for selected molecule
Ctrl+T	Project notes

Shortcut	Molecule operation
Ctrl+Alt+C	Copy 2D
Ctrl+C	Copy 3D
Ctrl+V	Paste
Ctrl+Space	Clear favorites

Shortcut	View/Display operation
Ctrl+Shift+R	Reset display
Ctrl+Alt+R	Rock display
Ctrl+Alt+S	Spin display
F11	Full screen
Shift+N	Reset labels

Right click to access menu
Atom
Molecule in table
Toolbars and window headers
Column title