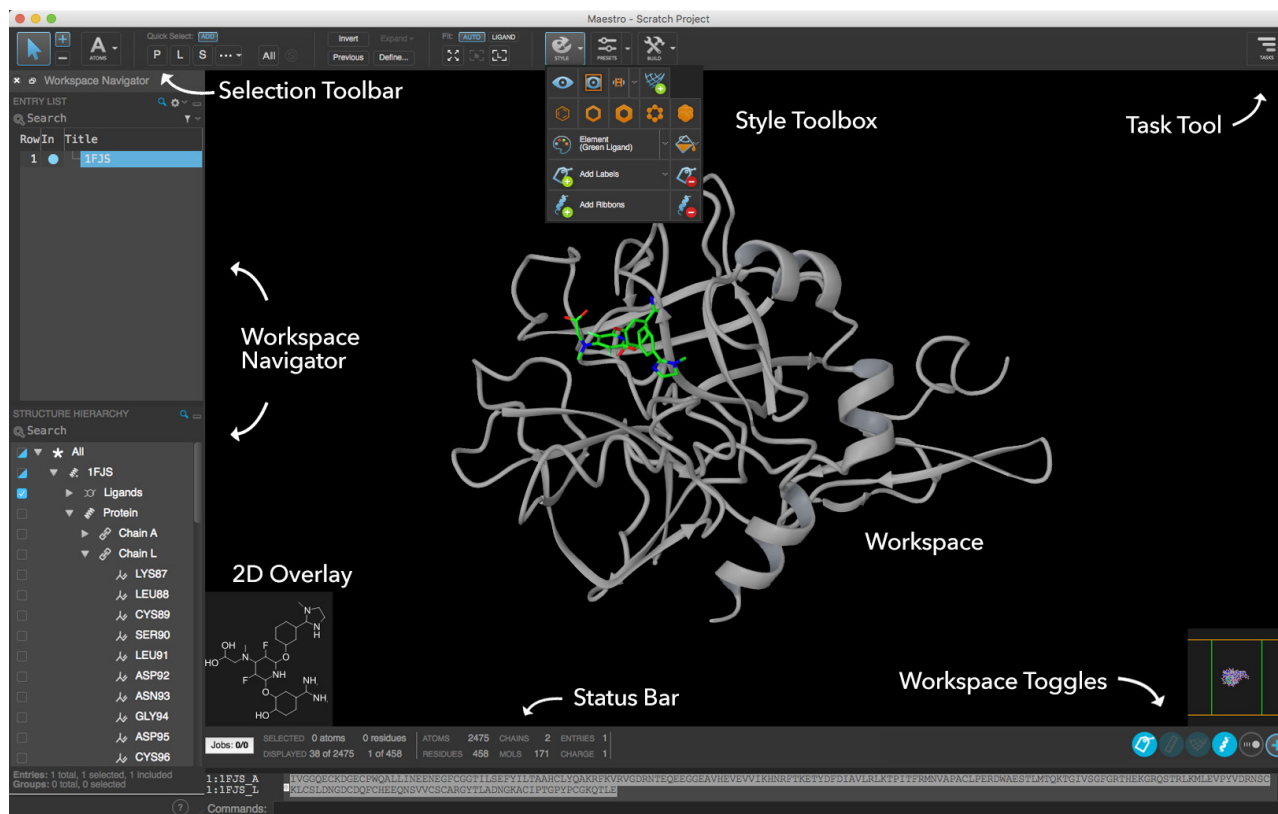


# Getting Started in Maestro 11



## Viewing Structures

### Mouse Controls

**Zoom:** Middle+right button or mouse scroll wheel

**Rotate xy:** Hold down middle button

**Rotate z:** Ctrl+ hold down middle button of mouse

**Translate xy:** Right button

**More mouse controls:** Maestro > Mouse Actions to view and modify additional controls.

**Select higher-level structure:** Double-click left button

**Stereo view:** Turn stereo view on and off using the panel opened from the Workspace Toggles bar, or type **Ctrl+S**.

**Clipping Planes:** Show or hide the gadget using the Workspace Toggles panel or the menu: Window > Clipping Planes. Click and drag the horizontal orange lines in the box to adjust the clipping planes' location.

**Protein Sequence:** Show or hide the gadget using the Workspace Toggles panel or the menu: Window > Sequence Viewer.

## Importing Structures

**Import your own files:** File > Import Structures; **Ctrl+I**


**Import a PDB file:** File > Get PDB

## Workspace Navigator


The Workspace Navigator includes the **Entry List** and the **Structure Hierarchy**. Each section can be resized or collapsed. Show or hide the whole panel using **Ctrl+E** or Window > Workspace Navigator.

**Select entries:** Click on the entry title in the Entry List. Selected rows are highlighted in blue.

**Include entries:** Toggle the circle next to the entry title. **Shift+Click** to include a range of entries in the Workspace. **Ctrl+Click** to add or remove an entry.

Use the **Structure Hierarchy** to locate and manipulate individual chemical components within each included entry. Click object names to select the corresponding atoms. Toggle visibility or hover and click  to style.


## Task Tool

Click the  button in the upper righthand corner to select from many common tasks. You can also begin typing a task keyword into the search field that appears (e.g., "docking"). Search results are dynamically updated.

**Hover** over an item in the Task Tool to display a description or **click** to open the application panel.

## Workspace Toggles


The Workspace Toggles bar contains several buttons for showing and hiding visual components of the Workspace. **Hover** over any button to see a description of what it does.

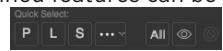
Click  to open a panel with additional Workspace configuration options.

## Selection Toolbar

You can quickly select or deselect chemical objects in the Workspace using the Selection Toolbar.


Click  or  to build a selection or remove items from it.

Change the picking level by clicking .

Several predefined features can be selected using the Quick Select buttons 

## Style Toolbox

Use the Style Toolbox to modify the visual representation of selected atoms in the Workspace.

Click  to modify the visual aspects of selected atoms.

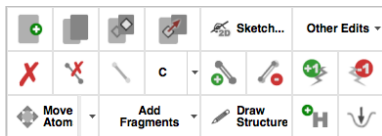
Click  for one-click styling of Workspace entries.



# Performing Common Tasks in Maestro 11

## Building and Modifying Structures


Click  to open the 3D Builder palette.

The 3D Builder palette can be used to edit existing entries or create new structures in the Workspace.






**2D building:** Click  Sketch... in the 3D Builder palette or select Edit > 2D Sketcher to open the 2D Sketcher. (This option is disabled when a macromolecule is in the Workspace.) To create a new, blank entry click the  button.


**3D building:** The 3D Builder palette contains many useful tools for modifying structures in the Workspace. Hover over any button to see a description. Add structural fragments by clicking the **Add Fragments** button or choose from several advanced modifications in the **Other Edits** menu. (Note: Draw Structure is temporarily disabled.)

**Add hydrogens:** Click  H in the 3D Builder palette to add missing hydrogens to all selected structures in the Workspace.

**Adjust torsions:** Select Edit > Adjust > Quick Torsion from the top menu, then click on a bond to select it. Click and drag left or right to adjust the torsion.

**Minimize structures:** Press **Ctrl+M** or click  to perform a quick minimization on selected structures in the Workspace. Press **Shift+Ctrl+M** to minimize only the ligand.



**Create new entries in Workspace:** The 3D Builder palette contains several options for creating new entries. Click  to create a new blank entry, or  to duplicate the currently included entry or entries.



**Measurements:** Select Workspace > Measure from the top menu. A banner will appear with a choice of measurement types (default is Distance). Select the appropriate atoms in the Workspace, and the value will appear. Click  in the Workspace Toggles bar to quickly hide/show measurements that have been previously created.

## Working with Multiple Structures


**Import single or multiple structures:** **Ctrl+I** or select File > Import Structures from the top menu.

**Clear the Workspace:** Select Workspace > Clear Entries to clear all entries from the Workspace.

**Tile entries:** When more than one entry is included in the Workspace, press **Ctrl+L** or click the  button in the Workspace Toggles bar and then click  in the panel that appears.

**Move one entry while keeping others fixed:** Click the  button in the Workspace Toggle bar and then toggle the  button on/off to choose whether tiles are controlled separately or together.

## Working with Proteins

Click  to open the Task Tool, then select Browse > Biologics to select from many protein-specific tasks.

**Generate a 2D diagram of a ligand in a binding pocket:** With a complex in the Workspace, select Browse > Structure Building > Ligand Interaction Diagram from the Task Tool to open the 2D Workspace. Here, select View > LID Legend for a diagram legend.

**Analyze protein quality and view Ramachandran plot:** Select Browse > Biologics > Structure Quality from the Task Tool to view a Ramachandran plot and Protein Report of potential problems with the structure in the Workspace.

## Getting Help


**Online Documentation:** Click  in any panel, or go to Help > Help...

**Tutorials:** Help > Tutorials...

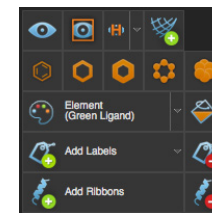
**Training Videos:** Visit [www.schrodinger.com/videos](http://www.schrodinger.com/videos)



**Knowledge Base:** Visit [www.schrodinger.com/kb](http://www.schrodinger.com/kb)

## Working with Styles

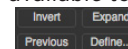
Click  to open the Style Toolbox.

The Style Toolbox provides you with precise control over how structures are displayed in the Workspace.



**Selectively display atoms:** Click the  button in the Style Toolbox to only display currently selected atoms or click the  button to display/undisplay currently selected atoms without affecting other atoms in the Workspace.

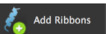

**Advanced selection options:** Several advanced selection modifiers are available to the right of the Quick Select buttons:




Use these options to invert the current selection, expand it by a given distance, or other property, or create complex selection definitions.

**Change molecular rendering:** Select the structures to be modified and then choose from several representations available in the Style Toolbox. Structures can be rendered in:

Wireframe , thin tubes , thick tubes , ball-and-stick , or CPK 

**Display ribbons:** Click  on the Style Toolbox to display protein ribbons for selected atoms. Click the  button in the Workspace Toggles bar to hide/show existing protein ribbons.

**Apply a predefined style:** Click  to choose from several predefined styles or to create your own.

**Create surfaces:** Click  in the Style Toolbox to quickly create a molecular surface on selected atoms. For more advanced options, use the Workspace > Surface menu.

# Keyboard Shortcuts in Maestro 11

## Project Operations

Show Project Table: **Ctrl+T**

New project: **Ctrl+N**

Open project: **Ctrl+O**

Close project: **Ctrl+W**

Import structures: **Ctrl+I**

Cut: **Ctrl+X**

Copy: **Ctrl+C**

Paste: **Ctrl+V**

Paste by placing: **Ctrl+Shift+V**, then click in the Workspace to place

Open Command Script Editor: **Ctrl+D**

Create Project Table entry from contents of Workspace: **Ctrl+Shift+N**

Display single-entry feedback in Workspace: **S**

Quit Maestro: **Ctrl+Q**

Open help page for active panel: **F1**

## Modifying Structures

Delete selected atoms: **Del**

Minimize selected atoms in the Workspace: Press **Ctrl+M** (minimizes currently selected atoms, or all atoms if none are selected)

Minimize ligand: **Shift+Ctrl+M** (minimizes all ligands currently included in the Workspace)

## Project Table Operations

Show Project Table: **Ctrl+T**

Mark all included entries in Project Table: **X**

Scroll up/down: **Up/down arrow keys**

Scroll up/down one page: **Page Up / Page Down**

Jump to top/bottom of the Project Table: **Home/End**

Jump to previous/next included entry: **Ctrl+Page Up/Down**

Jump to previous/next selected entry: **Shift+Page Up/Down**

Include the next or previous selected entry in the Workspace: **Right arrow** or **left arrow**, respectively

Display entry information in the Workspace: Press **S**

Include only selected entries in Workspace: **Ctrl+N** while the mouse pointer is over the Project Table

Exclude selected entries from Workspace: **Ctrl+X** while the mouse pointer is over the Project Table

## Finding and Selecting Atoms

Select single atom or bond with click in Workspace: **A**

Select residue, chain, molecule, or entry with click in Workspace: **R, C, M,** or **E**, respectively

Select all: **Ctrl+A**

Clear selection: **Ctrl+U**

Find substructures or entries: **Ctrl+F**

Select higher-level structure: **Double-click left button** on atom in Workspace

Select chemical object: **Click left button** on object in Structure Hierarchy

Select and fit to chemical object: **Double-click left button** on object in Structure Hierarchy

## Workspace Operations

Full screen Workspace mode: Press **Ctrl+=**; press **Esc** or **Ctrl+=** to exit

Apply Workspace style: **Ctrl+Y**

Tile Workspace: **Ctrl+L**

Fit Workspace to ligand: **L**

Fit to selected atoms: Press **Z**; if no atoms are selected, all Workspace contents will be fit to screen

Zoom in: **K**

Zoom out: **J**

Move clipping planes back: **-**

Move clipping planes forward: **+**

Move clipping planes together: **F**

Move clipping planes apart: **G**

Stereo view: **Ctrl+S**

Go to previous/next scene: Enter Scenes Mode and use **Ctrl+<** and **Ctrl+>**

## Using Saved Selections

Save selected atoms as Selection 1: Press **Ctrl+1** to store current selection; **Ctrl+0** through **Ctrl+9** work similarly

Use saved Selection 1 to select atoms: Press **1**; keys **0** through **9** work similarly