PyMOL by Schrödinger   Menu ≡
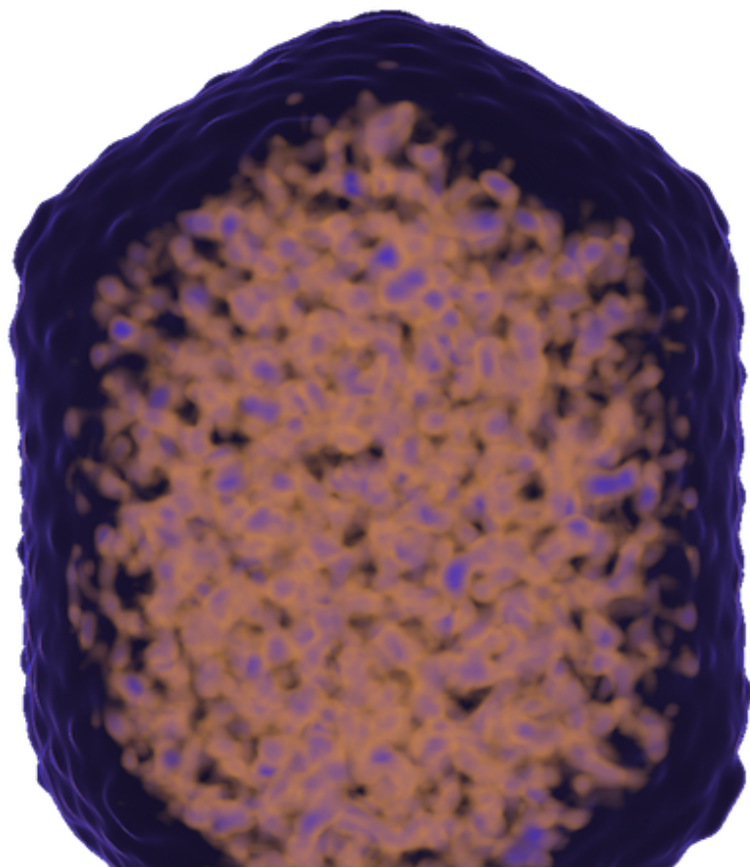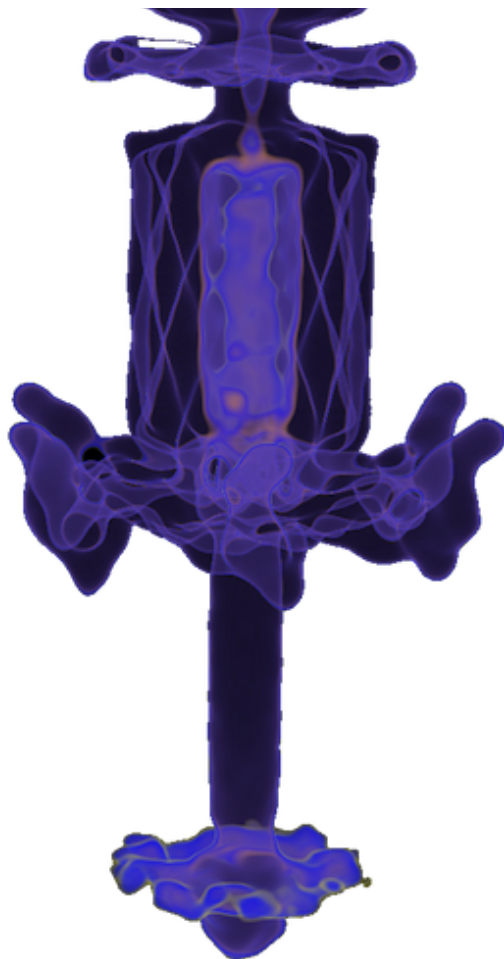
PyMOL is a user-sponsored molecular visualization system on an **open-source foundation**, maintained and distributed by Schrödinger.

We are happy to introduce
PyMOL 2.5!!

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RELEASE HIGHLIGHTS

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Version 2.5.2 - Updated August 20th 2021 (Installation instructions)

For previous versions, see here.

These bundles include Python 3.7.

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Windows

EXE Installer

Windows

ZIP Archive

macOS

DMG Disk Image

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TAR.BZ2 Archive

Or install from the Schrodinger Anaconda Channel.

`conda install -c schrodinger pymol-bundle`

New users:

[BUY LICENSE]

Existing users:

[DOWNLOAD LICENSE FILE]

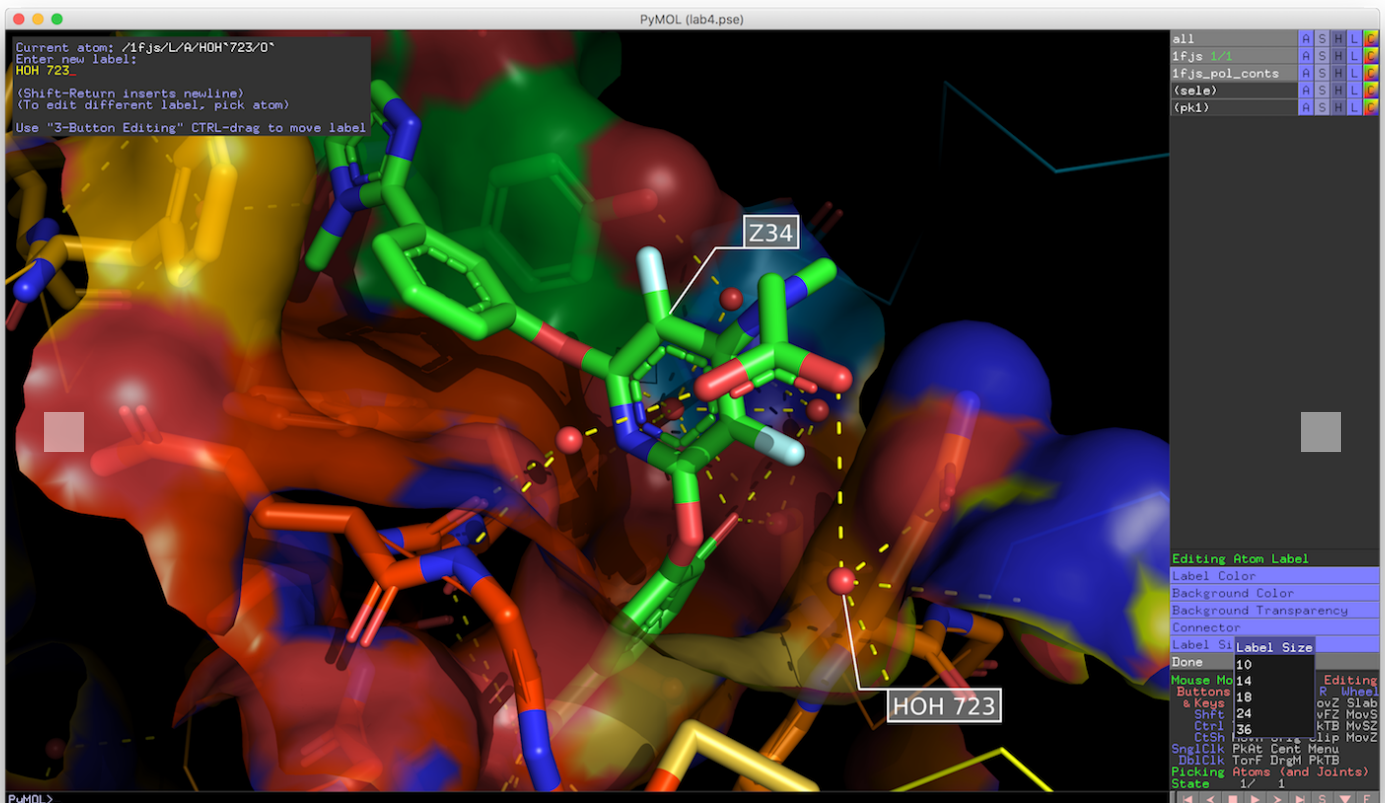# Release Highlights

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Unified modern user interface

PyQt interface replaces Tcl/Tk and MacPyMOL on all platforms

Anaconda Python distribution

Better third-party plugin and custom scripting support

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Liberal evaluation policy

Click here for a complete list of new features in PyMOL 2.5 (2.4) (2.3) (2.2) (2.1) (2.0)
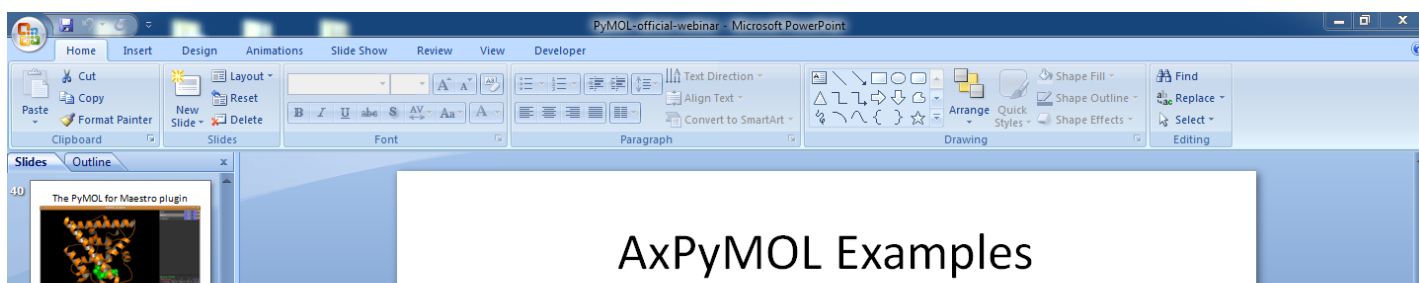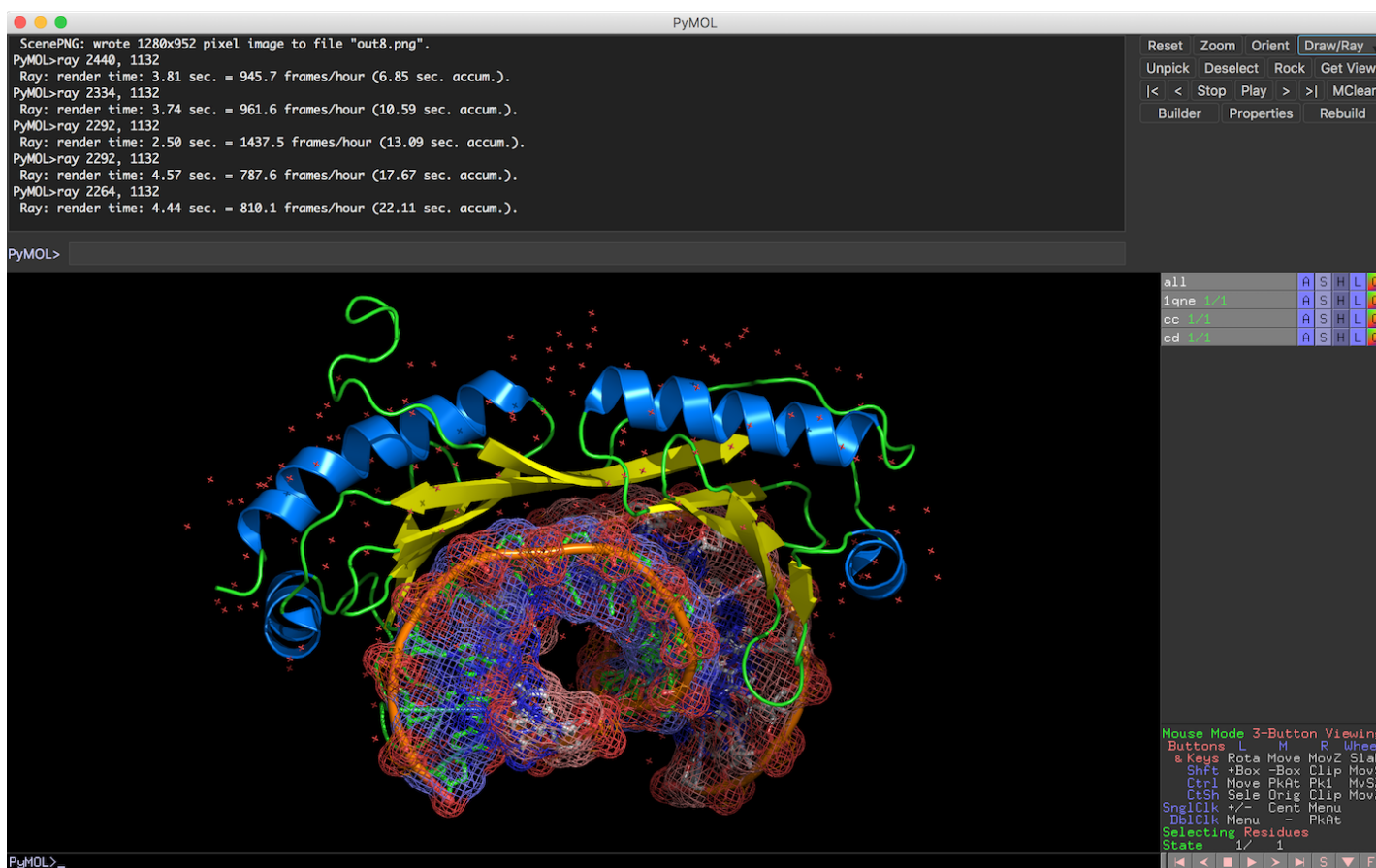
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Desktop PyMOL

A comprehensive software package for rendering and animating 3D structures

AxPyMOL

A plug-in for embedding 3D images and animations into PowerPoint presentations

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COMPARE PYMOL PRODUCTS

Open-Source Philosophy

PyMOL is a commercial product, but we make most of its source code freely available under a permissive license. The open source project is maintained by Schrödinger and ultimately funded by everyone who purchases a PyMOL license.

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DeLano.

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