

## Foldit Standalone

### Foldit Standalone is only licensed to entities that have a Rosetta™ license

The Foldit Standalone software provides a natural, intuitive interface to the [Rosetta](#) molecular modeling software. Foldit Standalone allows for user-centered protein manipulation for the purpose of solving structure prediction and design problems for proteins, and more generally other 3D structure related biochemistry problems. As an add-on to Rosetta, Foldit Standalone provides a complete novice and expert user interface for biochemistry structure exploration.

You can get a license to download Rosetta here: [Rosetta](#).

For more information: <http://fold.it/portal/>

For Descriptive and Technical Information: Seth Cooper [scooper@cs.washington.edu](mailto:scooper@cs.washington.edu)

### International Requests:

University of Washington agreements require additional review if the requesting entity is located in, or affiliated with the government of, China (Including Hong Kong; not including Taiwan), Iran, North Korea, Russia, or Syria. For requests from these countries, please **allow for an additional month of processing time** for a response.

### Available Licenses

#### Academic License:

See the panel to the right to license this product.

### Technology ID

45210

### Category

Software/Bioinformatics

Express Digital Content Licensing

### Learn more

