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## Welcome to:



*cisTEM* is user-friendly software to process cryo-EM images of macromolecular complexes and obtain high-resolution 3D reconstructions from them. It was originally developed by Tim Grant, Alexis Rohou and Nikolaus Grigorieff and comprises a number of tools to process image data including movies, micrographs and stacks of single-particle images, implementing a complete “pipeline” of processing steps to obtain high-resolution single-particle reconstructions. *cisTEM* is distributed under the [Janelia Research Campus Software License](#) and can be downloaded [here](#). We recommend downloading and using the pre-compiled binaries, rather than compiling the source code, for best performance. New users are encouraged to follow the [tutorial](#), which provides a quick way to become familiar with the most important functions of *cisTEM*.

### Support

Please visit the [documentation pages](#) for help. If you encounter any problems using *cisTEM*, or have questions, please use the [forums](#).

### S3 File:

[cistem-1.0.0-beta-intel-linux.tar.gz \(recommended\)](#) [cistem-1.0.0-beta-source-code.tar.gz](#)

[cisTEM Forum](#)

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