

Some softwares: (for single particle cryo-EM image processing)

- Imagic (van Heel lab) <https://www.imagescience.de/index.html>
- Spider (Frank lab) <https://spider.wadsworth.org>
- FREALIGN (Grigorieff lab) <http://grigoriefflab.janelia.org/frealign>
- EMAN (Ludtke lab) <http://blake.bcm.tmc.edu/EMAN2>
- XMIPP (Carazo lab) <http://xmipp.cnb.csic.es>
- Relion (Scheres lab) <http://www2.mrc-lmb.cam.ac.uk/relion>
- Scipion (Carazo lab) <http://scipion.cnb.csic.es/m/home>
- Sphire (Raunser lab) <http://sphire.mpg.de/>
- cisTEM (Grigorieff lab) <https://cistem.org>
- COW (Stark lab) www.cow-em.de
- cryoSPARC(2) (Punjani/Brubaker lab) <https://cryosparc.com/>
- Warp (Tegunov lab) <http://warpem.com>
- ...

Plus other specific software for helical reconstructions, viruses etc.

IHRSR (Egelman lab), SPRING (Sachse lab), Relion (Scheres lab)...
Baker lab, Zhou lab, etc.