

Project Help

SCIPION v3.0.0

View Protocols SPA

- Protocols SPA
 - Imports
 - Movies
 - Micrographs
 - CTF estimation
 - Preprocess
 - Particles
 - Picking
 - Extract
 - Preprocess
 - Filter
 - Mask
 - 2D
 - Align
 - Classifu
 - 3D
 - Initial volume
 - xmipp3 - ransac
 - xmipp3 - reconstruct.sier
 - xmipp3 - swarm consensus**
 - xmipp3 - convert a PDB
 - relicon - 3D initial model
 - eman2 - initial model
 - eman2 - initial model SGD
 - cruosparc2 - initial model
 - more
 - Preprocess
 - Classifu
 - Refine
 - Postprocess
 - Analysis
 - Reconstruct
 - Tools
 - Sets
 - Calculators
 - Exports

Protocol Run: XmippProtReconstructSwarm

mipp Protocol: xmipp3 - swarm consensus finished [Cite](#) [Help](#)

Run

Run name: `xmipp3 - swarm consensus` [Edit](#) Comment: [Edit](#)

Run mode: ☒ Continue ☐ Restart [?](#) Host: `localhost` [?](#)

Parallel Threads: `1` MPI: `12` [?](#) Use queue? ☐ Yes ☒ No [?](#)

GPU IDs: ☒ Yes ☐ No `0` [?](#) Wait for: [?](#)

Expert Level: ☒ Normal ☐ Advanced

Input

Full-size Images: `xmipp3 - center particles.outputParticles` [Search](#) [Delete](#) [View](#)

Initial volumes: `xmipp3 - align volume.outputVolumes` [Search](#) [Delete](#) [View](#)

Radius of particle (px): `-1` [?](#)

Symmetry group: `0` [?](#)

Mask: [Search](#) [Delete](#) [View](#)

[Close](#) [Save](#) [Execute](#)

inputParticles (from xmipp3 - center particles -> outputParticles) [out.SetOfParticles (3491 items, 120 x 120, 1.24 Å/px)]

inputVolumes (from xmipp3 - align volume -> outputVolumes) [out.SetOfVolumes (13 items, 120 x 120 x 120, 1.24 Å/px)]

Output

→ xmipp3 - swarm consensus -> outputVolume Volume (120 x 120 x 120, 1.24 Å/px)

→ xmipp3 - swarm consensus -> outputVolumes SetOfVolumes (13 items, 40 x 40 x 40, 3.71 Å/px)

SUMMARY

Symmetry: 0

Number of iterations: 15

