

View Protocols SPA

- Protocols SPA
 - Imports
 - Movies
 - Micrographs
 - CTF estimation
 - Preprocess
 - Particles
 - Picking
 - xmipp3 - particle boxsize
 - xmipp3 - manual-picking (step 1)
 - xmipp3 - auto-picking (step 2)
 - xmipp3 - deep consensus picking
 - xmipp3 - deep micrograph cleaner
 - xmipp3 - assign tiltpairs
 - xmipp3 - picking consensus
 - xmipp3 - pick noise
 - xmipp3 - remove duplicates
 - sphere - cruolo picking
 - sphere - cruolo training
 - relion - auto-picking
 - relion - auto-picking LoG
 - gautomatch - auto-picking
 - eman2 - boxer
 - eman2 - boxer auto
 - eman2 - sparx gaussian picker
 - cistem - find particles
 - Extract
 - Preprocess
 - Filter
 - Mask
 - 2D
 - Align
 - Classify
 - 3D
 - Initial volume
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 - Classify
 - Refine
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 - Analysis
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Protocol Run: XmippProtDeepMicrographScreen

mipp Protocol: xmipp3 - deep micrograph cleaner finished [Cite](#) [Help](#)

Run

Run name [Edit](#) Comment [Edit](#)

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

FGPU IDs ☒ Yes ☐ No [?](#) Use queue? ☐ Yes ☒ No [?](#)

Wait for [?](#)

Expert Level ☒ Normal ☐ Advanced

Input

Input coordinates [Search](#) [Delete](#) [Eye](#) [?](#)

Micrographs source ☒ same as coordinates ☐ other [?](#)

Threshold [?](#)

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

Summary Methods Output Log

Input

inputCoordinates (from create subset 0.5 -> outputCoordinates [outputCoord SetOfCoordinates (4006 items, 300 x 300)

Output

xmipp3 - deep micrograph cleaner -> outputCoordinates Full SetOfCoordinates (4006 items, 300 x 300)

xmipp3 - deep micrograph cleaner -> outputCoordinates_Manual_097 SetOfCoordinates (3968 items, 300 x 300)

Scipion3

View: Tree

