

view Protocols SPA

Micrographs

xmipp3 - preprocess micrographs

CTF estimation

gctf - ctf estimation

relion - export ctf

xmipp3 - ctf estimation

xmipp3 - ctf consensus

eman2 - ctf auto

grigoriefflab - ctffind4

more

Particles

Picking

relion - auto-picking

xmipp3 - particle boxsize

xmipp3 - manual-picking (step 1)

xmipp3 - auto-picking (step 2)

xmipp3 - deep consensus picking

xmipp3 - assign tiltpairs

xmipp3 - picking consensus

sphire - cryolo picking

sphire - cryolo training

eman2 - boxer

eman2 - boxer auto

eman2 - sparx gaussian picker

bsoft - particle picking

Extract

Preprocess

Filter

xmipp3 - filter particles

xmipp3 - denoise particles

Protocol Run: ProtRelionAutopickLoG

RELION Protocol: relion - auto-picking LoG finished Cite Help

Run

Run name relion - auto-picking LoG Comment

Run mode ☒ Continue ☐ Restart Host localhost

Parallel ☐ Threads ☒ MPI 4 Use queue? ☐ Yes ☒ No

Wait for

Input Streaming

Input

Input micrographs xmipp3 - ctf consensus.outputMicrographs

Box size (px) 300

Laplacian of Gaussian

Diameter for LoG filter (A) Min 250 Max 300

Are the particles white? ☐ Yes ☒ No

Maximum resolution to consider (A) 20.0

Adjust default threshold 0.2

Additional arguments:

Close Save Execute

Input

inputMicrographs (from xmipp3 - ctf consensus -> outputMicrographs [outputMicrographs])

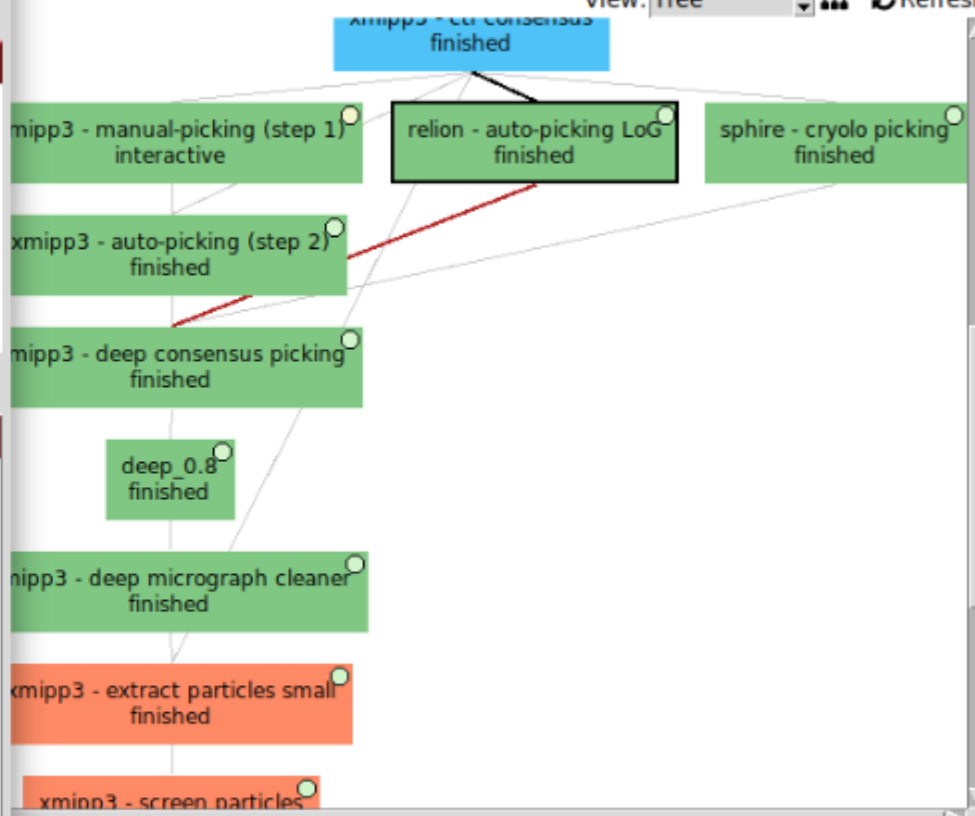
Output

relion - auto-picking LoG -> outputCoordinates

Sample_10248

Protocols | Data

View: Tree Refresh



Analyze Results

SetOfMicrographs (47 items, 3710 x 3838, 0.49 Å/px)

SetOfCoordinates (5813 items, 300 x 300)