


View Protocols SPA

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Protocol Run: XmippProtCenterParticles


Protocol: xmipp3 - center particles

finished
📄 Cite
❓ Help

Run

Run name xmipp3 - center particles ✎ Comment ✎

Run mode ☒ Continue ☐ Restart ❓ Host localhost ▼

Use queue? ☐ Yes ☒ No ✎ ❓

Wait for ❓

Input

Input Classes noFerritin classes.outputClasses2D 🔍 🗑️ 👁️ ❓

Set of micrographs xmipp3 - ctf consensus GCTF+CTFFind4+Xmipp.outputMicrographs 🔍 🗑️ 👁️ ❓

✕ Close
💾 Save
⚙️ Execute

```

▼ Input
  inputClasses (from noFerritin classes -> outputClasses2D [outputClasses: SetOfClasses2D (38 items)
  inputMics (from xmipp3 - ctf consensus GCTF+CTFFind4+Xmipp -> outputMicrographs (28 items, 3710 x 3838, 0.49 Å/px)
▼ Output
  ➡ xmipp3 - center particles -> outputClasses SetOfClasses2D (38 items)
  ➡ xmipp3 - center particles -> outputParticles SetOfParticles (3491 items, 120 x 120, 1.24 Å/px)
  ➡ xmipp3 - center particles -> outputCoordinates SetOfCoordinates (3491 items, 300 x 300)

SUMMARY
Realignment of 38 classes.
  
```

Protocol

View: Tree

finished

xmipp3 - center particles
finished- 3D initial model
finished

xmipp3 -

Analyze