

View Protocols SPA

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 **Protocol Run: XmippProtRansac** finished [Cite](#) [Help](#)

Run

Run name `xmipp3 - ransac`

Run mode ☒ Continue ☐ Restart

Parallel ☒ Threads ☐ MPI

GPU IDs ☒ Yes ☐ No

Expert Level ☒ Normal ☐ Advanced

Host `localhost`

Use queue? ☐ Yes ☒ No

Wait for

Input

Input averages `xmipp3 - center particles.outputClasses`

Symmetry group

Close Save Execute

Summary Methods Output Log

▼ Input

input.Set. (from xmipp3 - center particles -> outputClasses [output.Clas SetOfClasses2D (38 items)

▼ Output

xmipp3 - ransac -> outputVolumes SetOfVolumes (10 items, 120 x 120 x 120, 1.24 Å/px)

SUMMARY

RANSAC iterations: 400
Number of volumes to refine: 10
Number of refinement iterations: 10

View: Tree

osparc2 - initial model
finishedxmipp3 - ransac
finished3 - crop/resize volumes (2)
finishedxmipp3 - align volume
finished[Analyze Results](#)