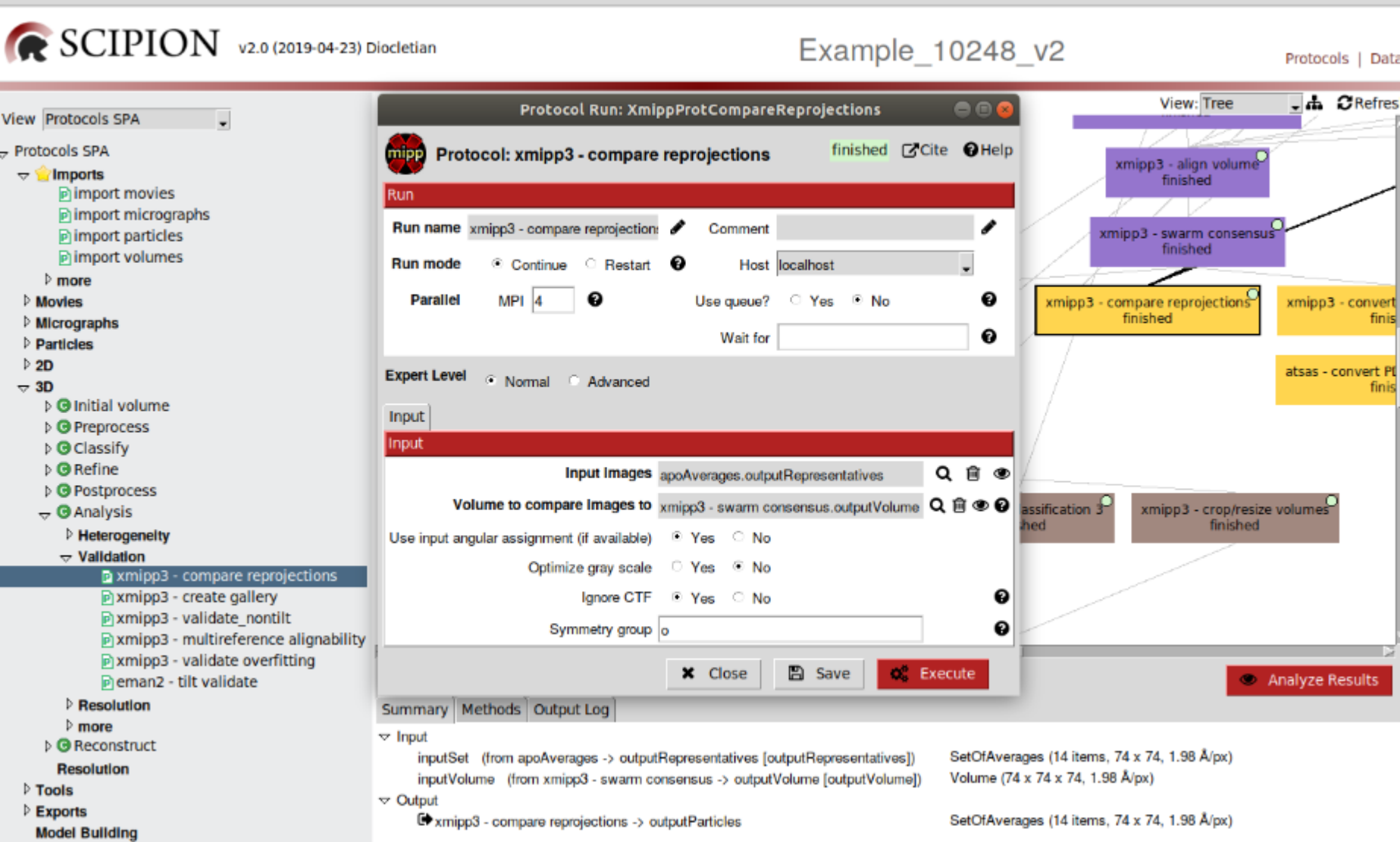
SCIPION
v2.0 (2019-04-23) Diocletian
Example_10248_v2
Protocols | Data
View Protocols SPA
Protocols SPA
Imports
import movies
import micrographs
import particles
import volumes
more
Movies
Micrographs
Particles
2D
3D
Initial volume
Preprocess
Classify
Refine
Postprocess
Analysis
Heterogeneity
Validation
xmipp3 - compare reprojections
xmipp3 - create gallery
xmipp3 - validate_nontilt
xmipp3 - multireference alignability
xmipp3 - validate overfitting
eman2 - tilt validate
Resolution
more
Reconstruct
Resolution
Tools
Exports
Model Building
Protocol Run: XmippProtCompareReprojections
Protocol: xmipp3 - compare reprojections
finished
Cite
Help
Run
Run name
xmipp3 - compare reprojection
Comment
Run mode
Continue
Restart
Host
localhost
Parallel
MPI
4
Use queue?
Yes
No
Wait for
Expert Level
Normal
Advanced
Input
Input
Input Images
apoAverages.outputRepresentatives
Volume to compare Images to
xmipp3 - swarm consensus.outputVolume
Use input angular assignment (if available)
Yes
No
Optimize gray scale
Yes
No
Ignore CTF
Yes
No
Symmetry group
o
Close
Save
Execute
View: Tree
Refres
xmipp3 - align volume
finished
xmipp3 - swarm consensus
finished
xmipp3 - compare reprojections
finished
xmipp3 - convert
finis
atsas - convert PD
finis
assification 3
hed
xmipp3 - crop/resize volumes
finished
Analyze Results
Summary
Methods
Output Log
Input
inputSet (from apoAverages -> outputRepresentatives [outputRepresentatives])
SetOfAverages (14 items, 74 x 74, 1.98 Å/px)
inputVolume (from xmipp3 - swarm consensus -> outputVolume [outputVolume])
Volume (74 x 74 x 74, 1.98 Å/px)
Output
xmipp3 - compare reprojections -> outputParticles
SetOfAverages (14 items, 74 x 74, 1.98 Å/px)