

View Protocols SPA

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 - eman2 - refine 2D bispec
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Protocol Run: ProtUserSubSet

Protocol: pwem - ProtUserSubSet finished [Cite](#) [Help](#)

Run

Run name Comment

Run mode ☒ Continue ☐ Restart Host

Use queue? ☐ Yes ☒ No

Wait for

Summary Methods Output Log

Input
inputObject (from crusparc2 - 2d classification)

Output
noFerritin classes -> outputClasses2D

SUMMARY

From input SetOfClasses2D of size 50 we selected 38 items to create output SetOfClasses2D

pion3

Prot

View: Tree

xmipp3 - extract particles
finishedcryosparc2 - 2d classification
finishednoFerritin classes
finishedxmipp3 - center particles
finishedrelion - 3D initial model
finished

xmipp3 -