

View **Protocols SPA**

Protocols SPA

- Imports
  - import movies
  - import micrographs
  - import particles
  - import volumes
- more
  - import coordinates
  - import ctfs
  - import atomic structure
  - import sequence
  - import averages
  - import masks
  - import micrograph pairs
  - export to EMDb**
- Movies
- Micrographs
  - xmipp3 - preprocess micrograph
  - xmipp3 - ctf estimation
  - CTF estimation
- more
- Particles
  - Picking
  - Extract
  - Preprocess
  - Filter
  - Mask

**Protocol Run: ProtExportEMDB**

**Protocol: scipion - export emdb** [Cite](#) [Help](#)

**Run**

Run name **scipion - export emdb (copy)** [Edit](#) Comment [Edit](#)

Run mode ☒ Continue ☐ Restart [?](#) Host **localhost**

Use queue? ☐ Yes ☒ No [?](#)

Wait for [?](#)

**Input**

**Input**

Volume to export	import half1.outputVolume.	<a href="#">Search</a> <a href="#">Delete</a> <a href="#">Eye</a> <a href="#">Help</a>
FSC to export	xmipp3 - resolution 3D	<a href="#">Search</a> <a href="#">Delete</a> <a href="#">Eye</a> <a href="#">Help</a>
Atomic structure to export	import atom struct mmCIF 5ni1.cif.outputPdb.	<a href="#">Search</a> <a href="#">Delete</a> <a href="#">Eye</a> <a href="#">Help</a>
Export to directory	/home/Submission1_03_24_2019	<a href="#">Folder</a> <a href="#">Help</a>

**Close** **Save** **Execute**