














View Protocols SPA

Protocols SPA

Imports



 import movies import micrographs import particles import volumes

more

 import coordinates import ctfs import atomic struc **import sequence** import averages import masks import micrograph export to EMX export to EMDB

Movies

Micrographs

 xmipp3 - preprocess m CTF estimation

Particles

 Picking Extract Preprocess Filter

Edit Copy

scipion - import

Summary Met

Input

Output

 scipion

SUMMARY

Sequence of aminoacids:

Sequence **HBA_HUMAN** imported from UniProt ID **P69905**

Protocol Run: ProtImportSequence



Protocol: scipion - import sequence

finished

 Cite Help

Run

Run name scipion - import sequence HB/

Comment

Run mode

☒ Continue☐ Restart

Host

localhost

Use queue?

☐ Yes☒ No

Wait for

Input

Input

Sequence ID

Sequence name

HBA_HUMAN

Sequence description

Import sequence of



☒ aminoacids☐ nucleotides

From

☐ plain text☐ atomic structure☐ file☒ UniProt ID

UniProt name/ID

P69905

 Close Save Execute Analyze Results