

View Protocols SPA

## Protocols SPA

## Imports

- import movies
- import micrographs
- import particles
- import volumes

## more

- import coordinates
- import ctfs
- import PDB
- 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



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

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Protocols | Da

Tree

Refresh

3



Analyze Results

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