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Protocol Run: ProtImportSequence

Protocol: scipion - import sequence finished [Cite](#) [Help](#)

**Run**

Run name **scipion - import sequence HBA** [Edit](#) Comment [Edit](#)

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

Use queue? ☐ Yes ☒ No [?](#)

Wait for [?](#)

**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](#)

**Summary** **Methods** **Output Log**

Input

Output

scipion - import sequence HBA\_HUMAN -> outputSequence

Sequence (name = HBA\_HUMAN)

**SUMMARY**

Sequence of aminoacids:

Sequence **HBA\_HUMAN** imported from UniProt ID **P69905**

[Analyze Results](#)