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### Protocol Run: ChimeraProtOperate

**Protocol: chimerax - operate** finished [Cite](#) [Help](#)

**Run**

Run name: **chimerax - operate** [Edit](#) Comment: [Edit](#)

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

Use queue? ☐ Yes ☒ No [Edit](#) [Help](#)

Wait for: [Edit](#)

[Input](#) [Help](#)

**Input**

Input Volume	Object	Info
Input additional Volumes		

**Atomic structure** **pwem - import atomic structure 5ni1.outputPdb** [Q](#) [C](#) [E](#) [I](#) [H](#)

Other atomic structures	Object	Info

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

**Input**

pdbFileToBeRefined (from pwem - import atomic structure 5ni1 -> outputPdb [out] AtomStruct (pseudoatoms=False, volume=False)

**Output**

[chimerax - operate -> 5ni1\\_chainA\\_Atom\\_struct\\_\\_3\\_007965](#) AtomStruct (pseudoatoms=False, volume=False)

**SUMMARY**

Produced files:  
we have some result