

Protocol: chimera - contacts

finished Cite Help

Run

Run name: chimera - contacts (unit cell) Comment

Run mode: ☒ Continue ☐ Restart Host: localhost

Use queue? ☐ Yes ☒ No

Wait for

Expert Level: ☒ Normal ☐ Advanced

Input

Atomic Structure: atomstructutils - operator (unit cell).outputPdb

Chain Labeling: {"A": "chainA", "A002": "HEM_A", "B": "chainB", "B002": "H

Apply symmetry: ☒ Yes ☐ No

Symmetry: Cn

Symmetry Order: 2

Close Save Execute

Entry Grid

	label
A	chainA
A002	HEM_A
B	chainB
B002	HEM_B

OK

