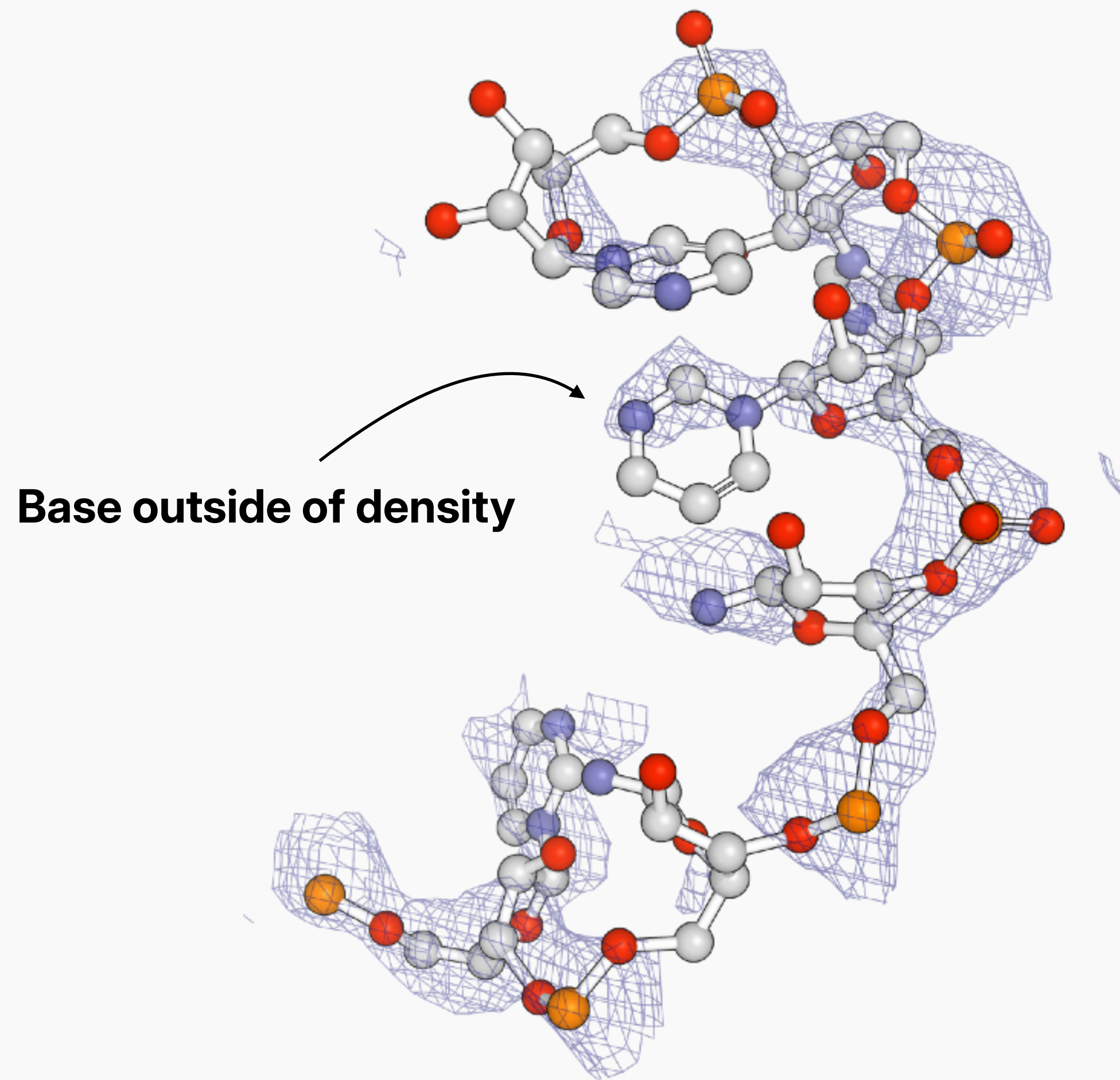


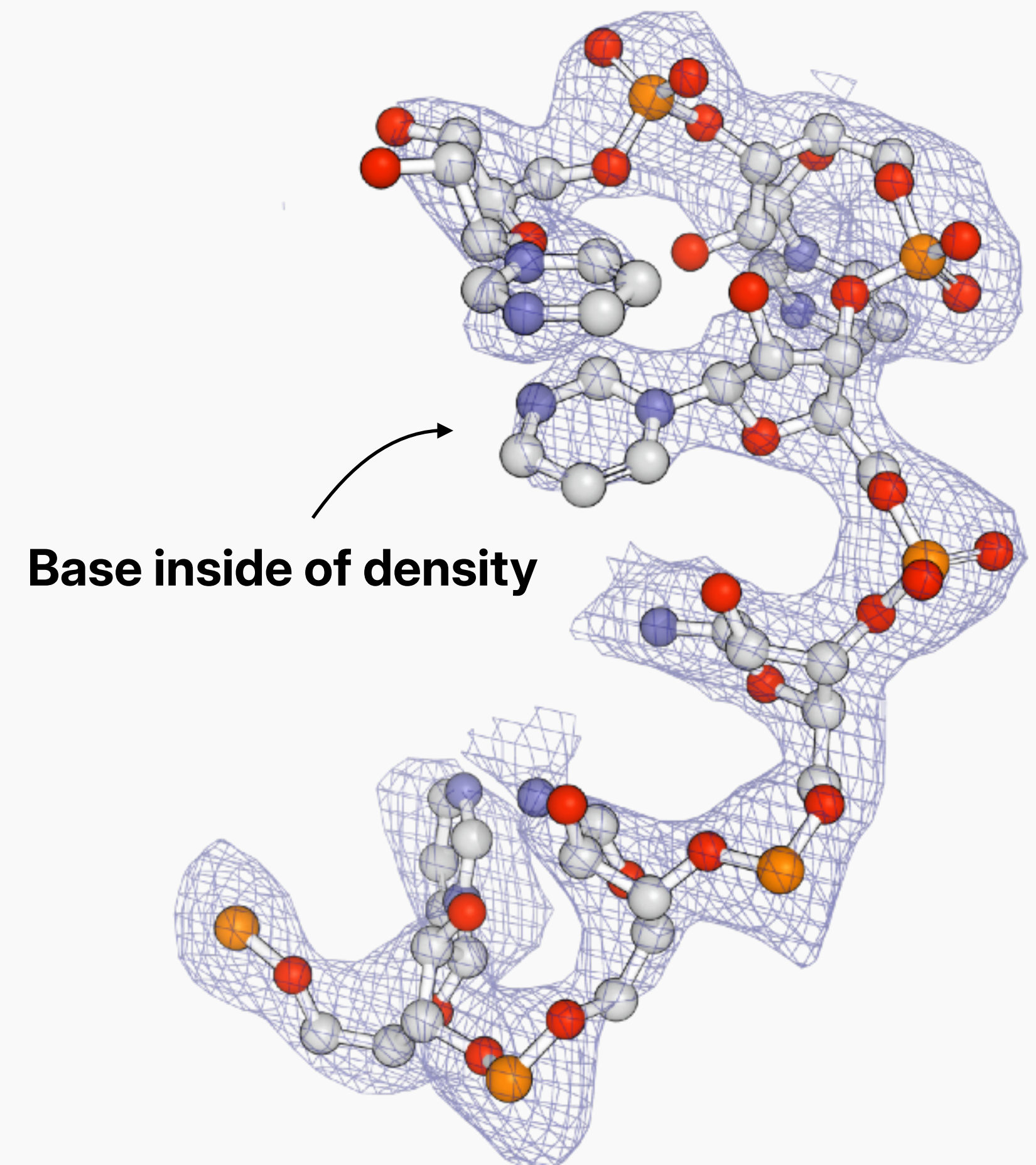
Nucleic Acid Structure Solution

Reciprocal Space Refinement



Servalcat

Default
Parameters



NucleoFind Result
PDB Code: 9LRI

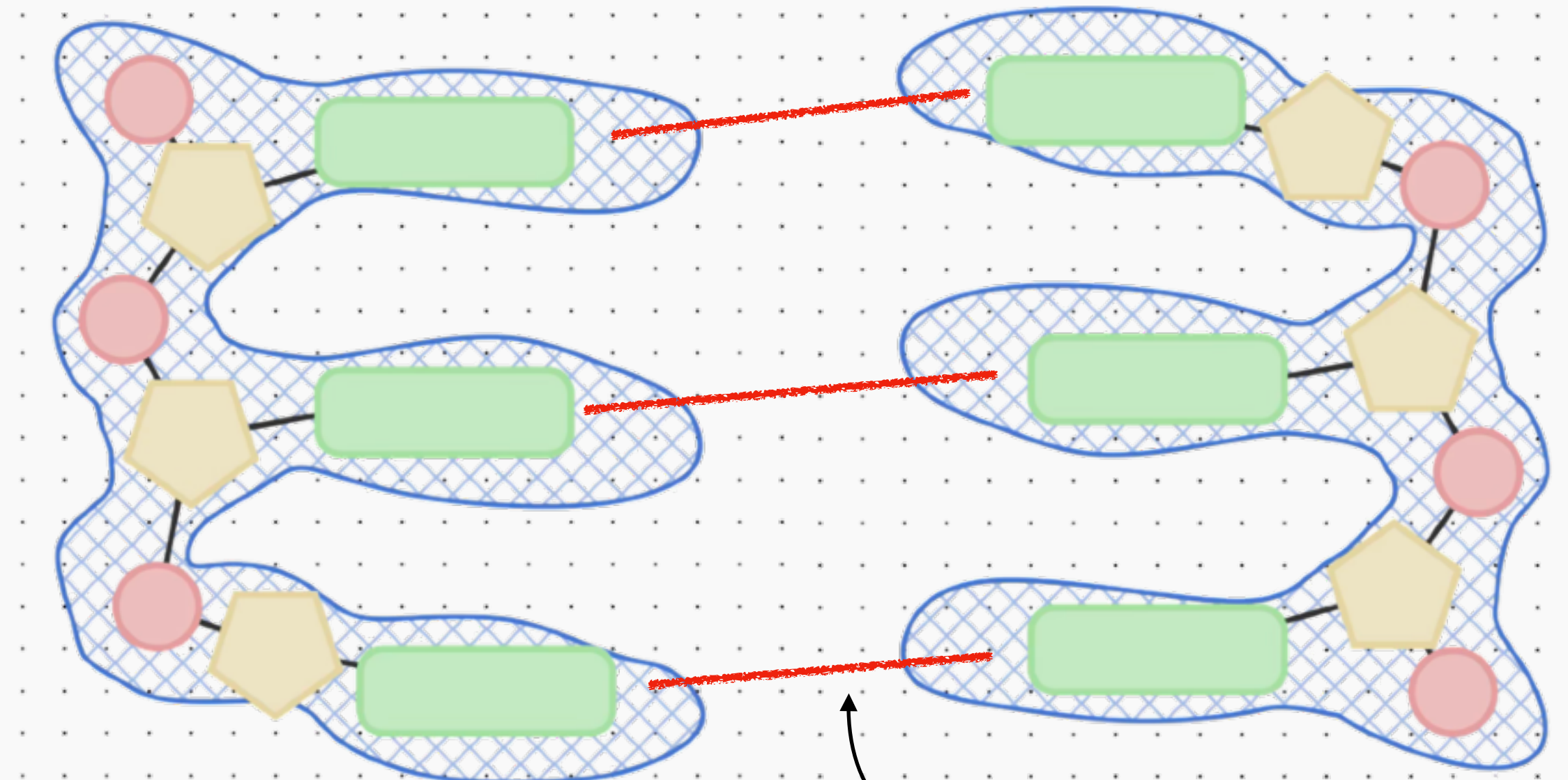
Nucleic Acid Structure Solution

Refinement Restraints

Base pairing and puckering restraints

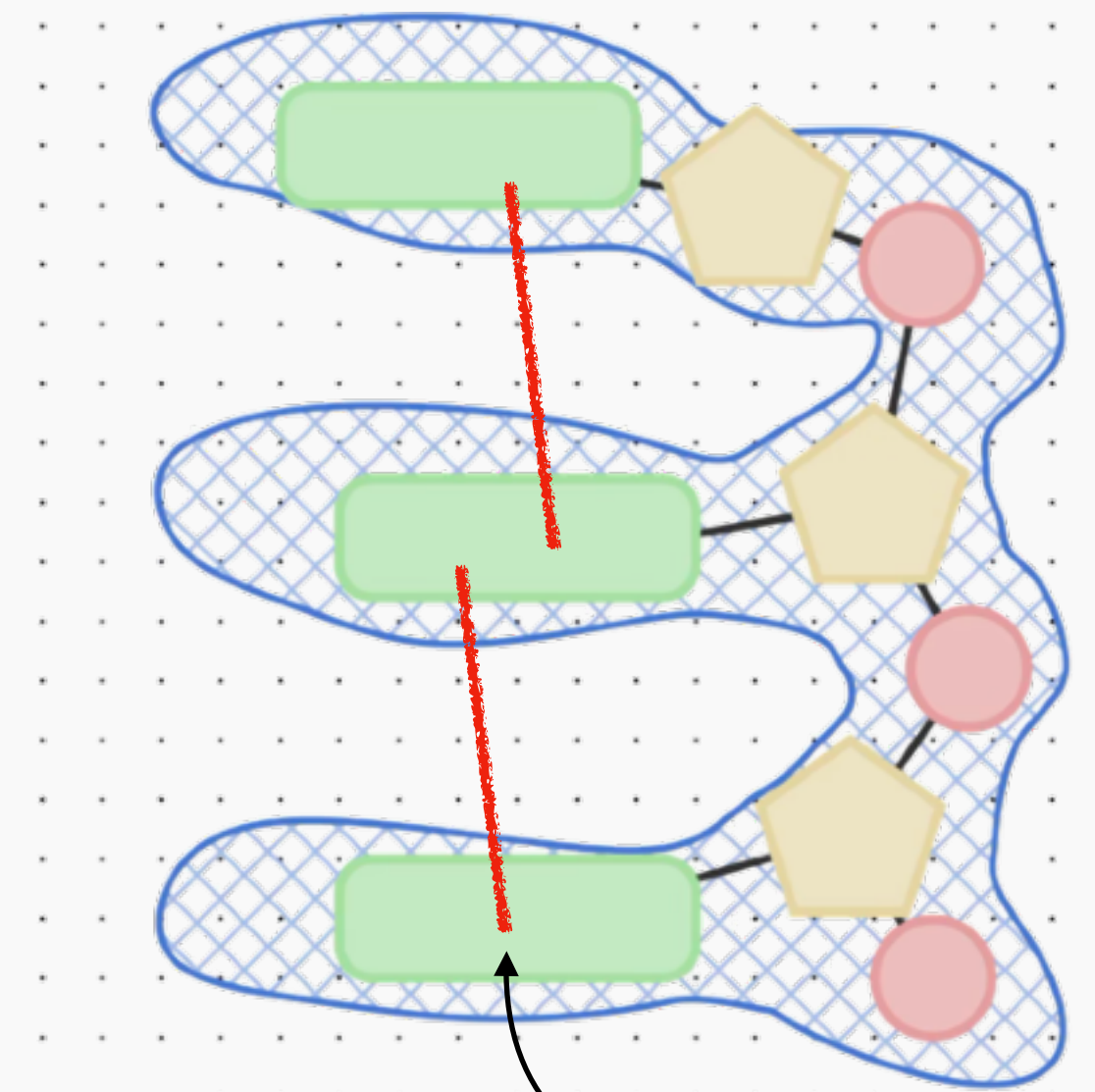
Libg

DoubleHelix



Base Pairing Restraint

Conformer restraints



Conformer Restraint