

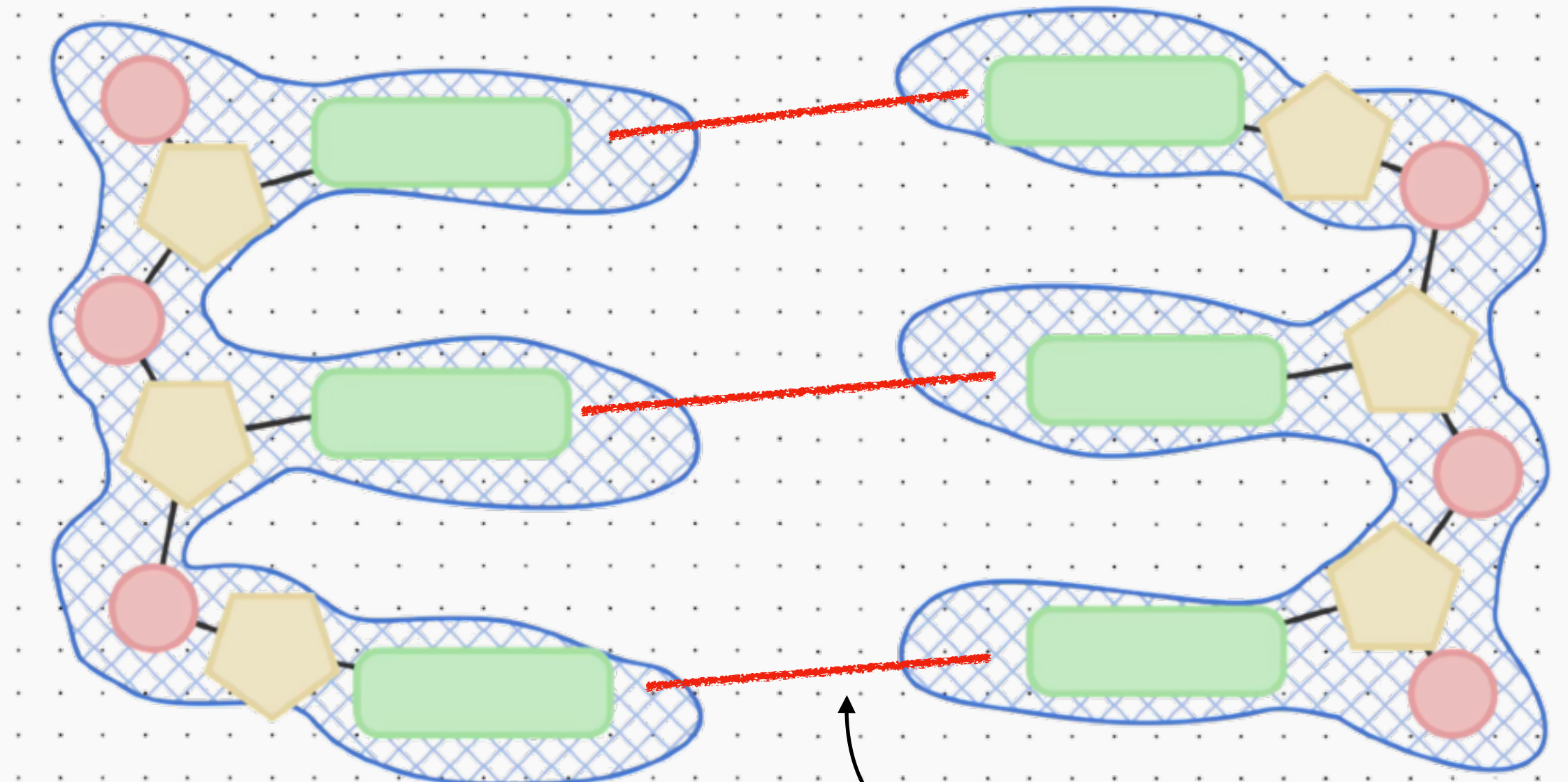
Nucleic Acid Structure Solution

Refinement Restraints

Base pairing and puckering restraints

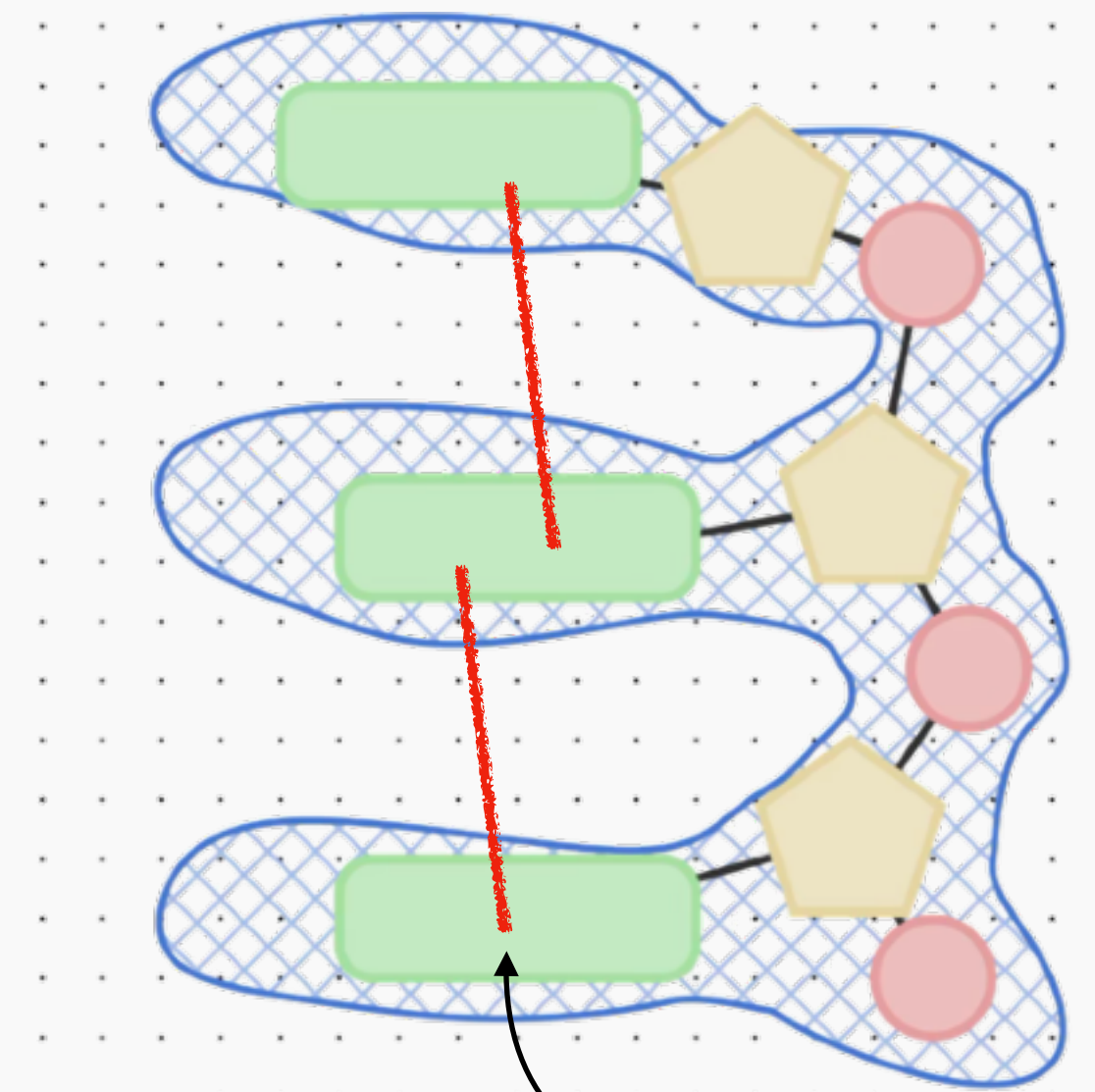
Libg

DoubleHelix



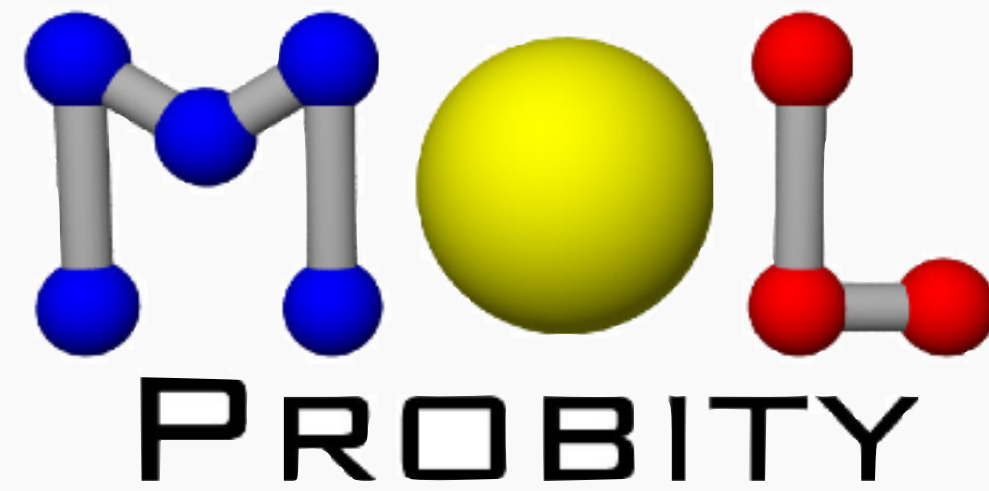
Base Pairing Restraint

Conformer restraints

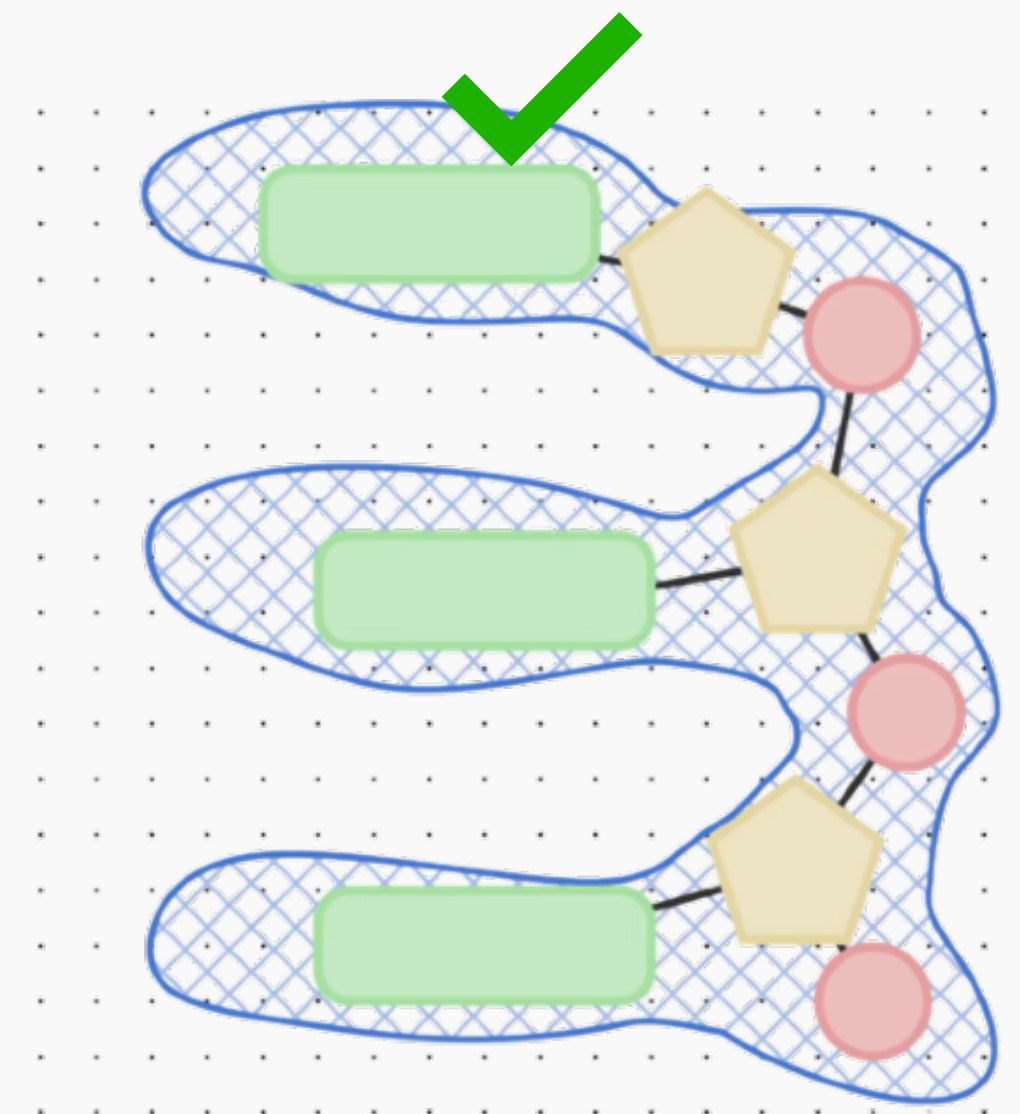
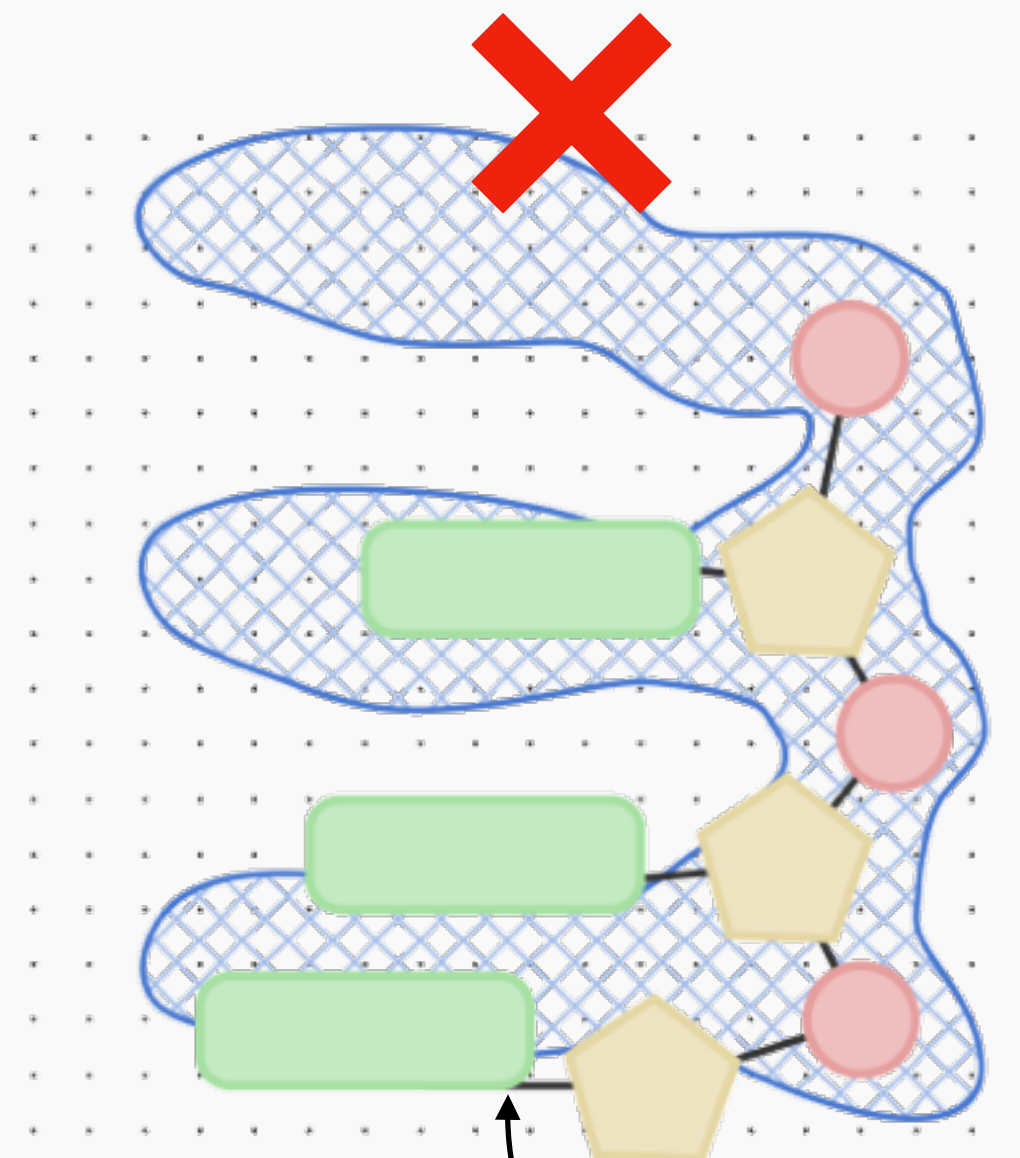


Conformer Restraint

Nucleic Acid Structure Solution Validation



DoubleHelix



Mismodelled