

X

1

1. Identify a region important for binding

2. Search for the best linkersize

3. Findlinker sequence



Background

Project Description

Results

Conclusion

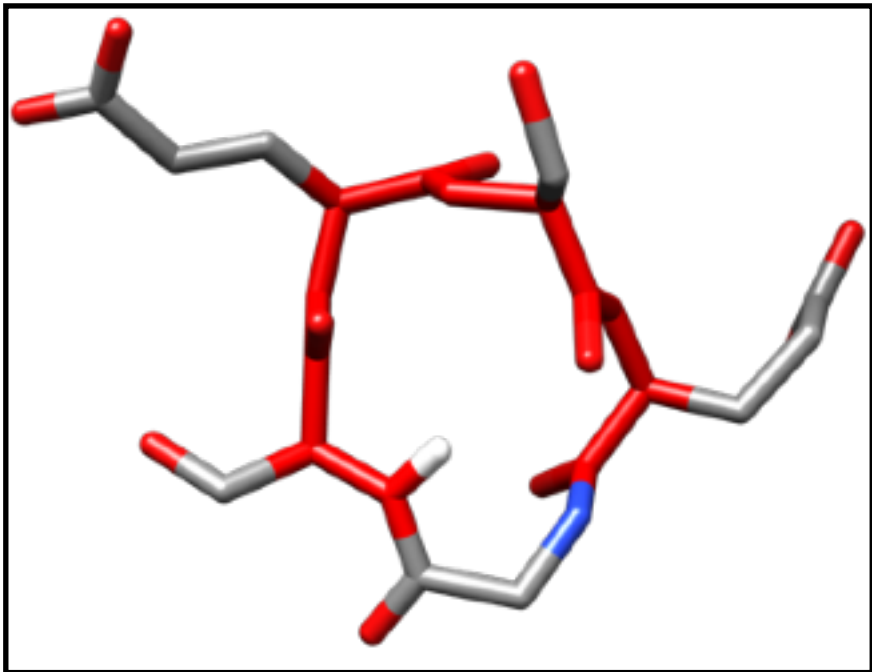
Cyclic-peptide design is a 3-step process

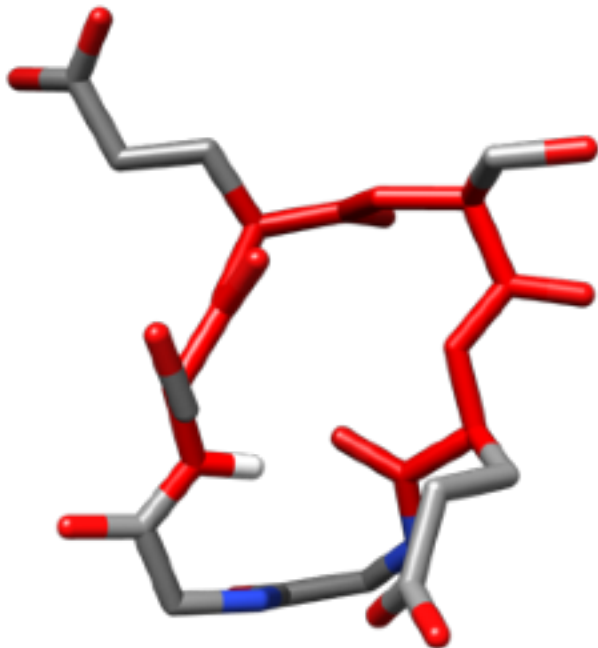
X₂

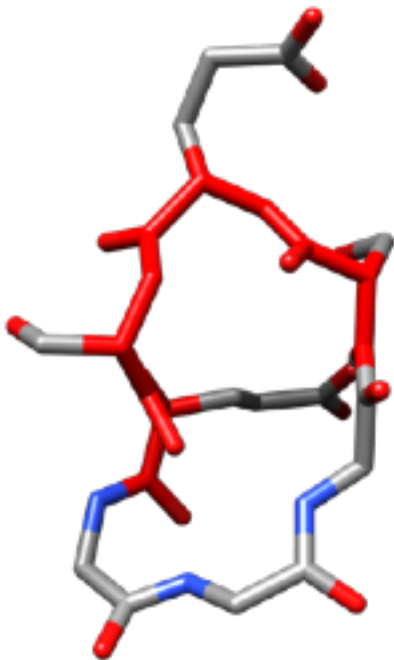


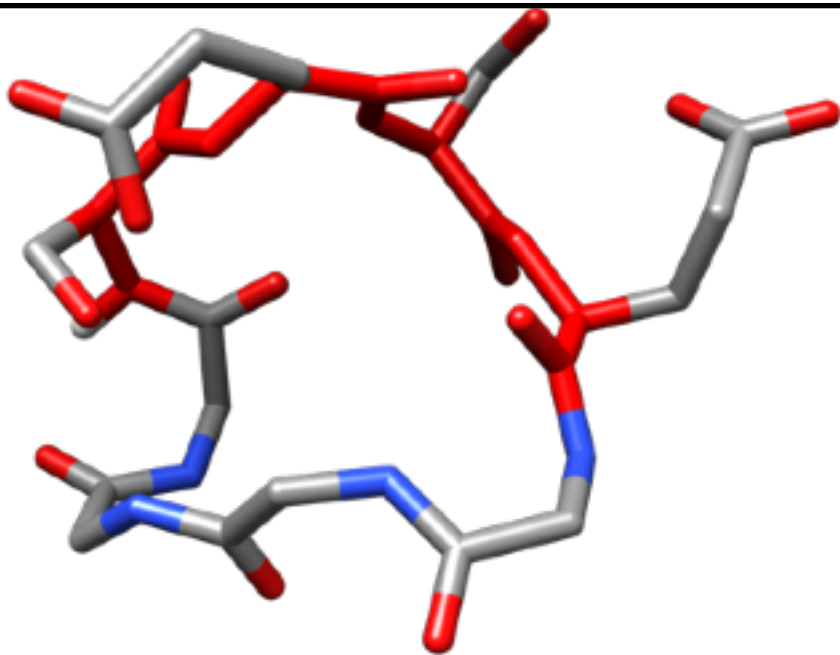
Xn

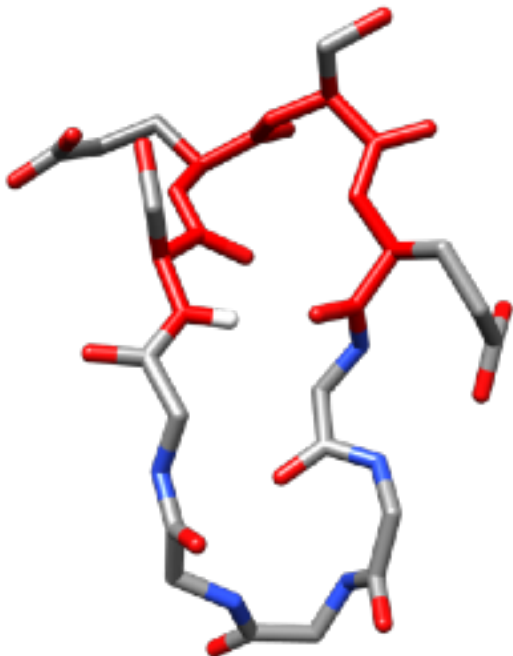
Create models and simulations

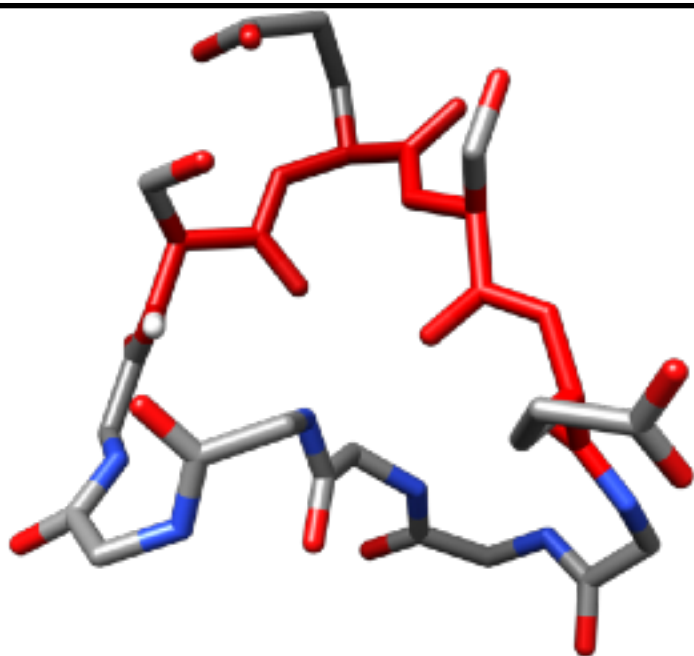












cyclo-SEES-SEC-Gr1

cyclo-SEESSECG2

cyclo-SEESSEEG3

cyclo-SEES-SEG4

cyclo-SEESSECG5

cyclo-SEESSEEG6