Deep Learning Approaches for the Design of Symmetric Cyclic Peptide Complexes





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Introduction

Background: Cyclic peptides have demonstrated promising potential for use as drugs and other molecular functionalities. However, the *de novo* design of such molecules has been mostly limited to classical, energybased design techniques and focused on the design of cyclic peptides as monomers rather than as homooligomer species.

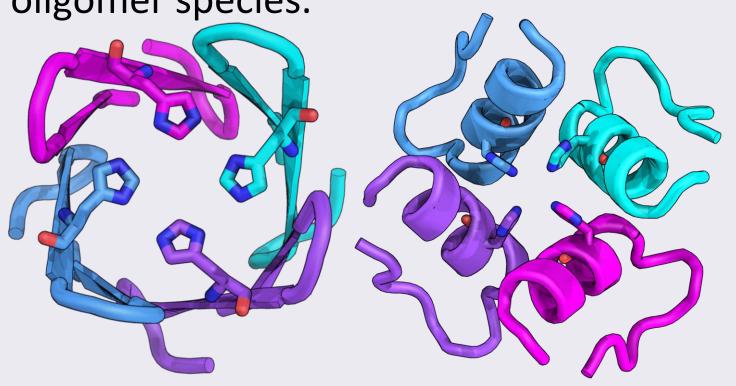


Figure 1. Cyclic peptide designs for nickel binding generated by Gaurav Bhardwaj.

Objective

- Explore the design of homooligomeric cyclic peptide systems using Deep learning.
- Design a protocol to generate cyclized peptide oligomers exhibiting "cage-like" symmetries in high-order complexes such as icosahedrons and octahedrons.

Design Pipeline

RFdiffusion¹: Protein structure generative diffusion model. **ProteinMPNN²:** Protein sequence design model using message-passing neural networks given backbone structures. AFCycDesign³: a variant of AlphaFold2 to predict the cyclic peptide structures.

Methods

Sample Dataset

- 4320 generated icosahedral oligomers of cyclic peptides using 4 different RFdiffusion design hyperparameters and 2 guiding potentials (e.g. inter/intra molecular weights).
- 59718 extracted C3 and C5 cyclic peptides from backbones of generated oligomers.

Design Protocol

- 1. Generate cyclized peptide oligomers of icosahedral symmetry using RFdiffusion.
- 2. Design symmetric sequences of those oligomers using ProteinMPNN.
- 3. Extract C3/C5 peptides from those sequence designs.
- 4. Predict and validate the accuracy of extracted C3/C5 peptides using AfCycDesign.

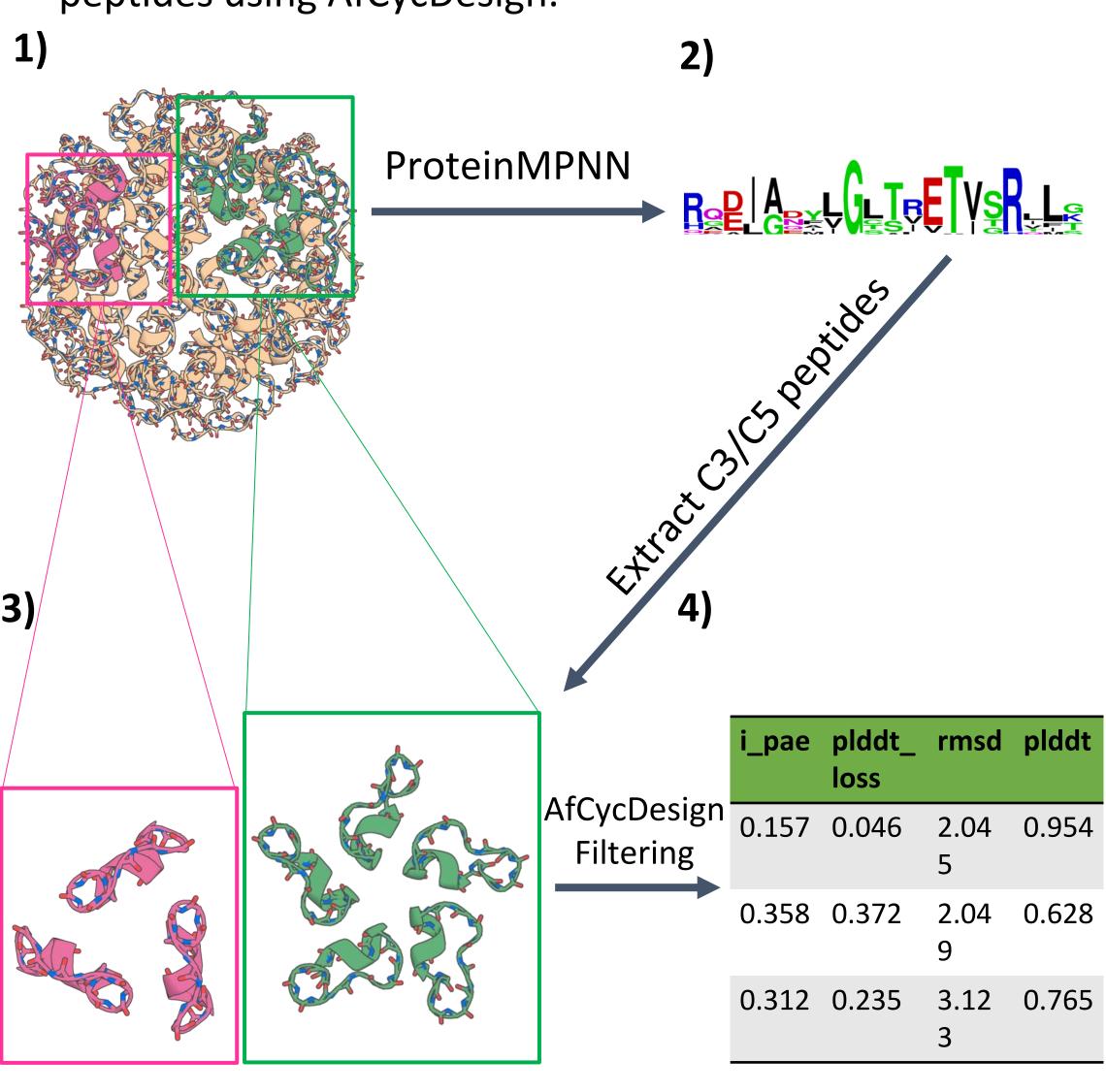


Figure 3. Design protocol to generate cyclic peptide oligomers exhibiting icosahedral symmetries.

Predict how symmetric

sequences will fold via

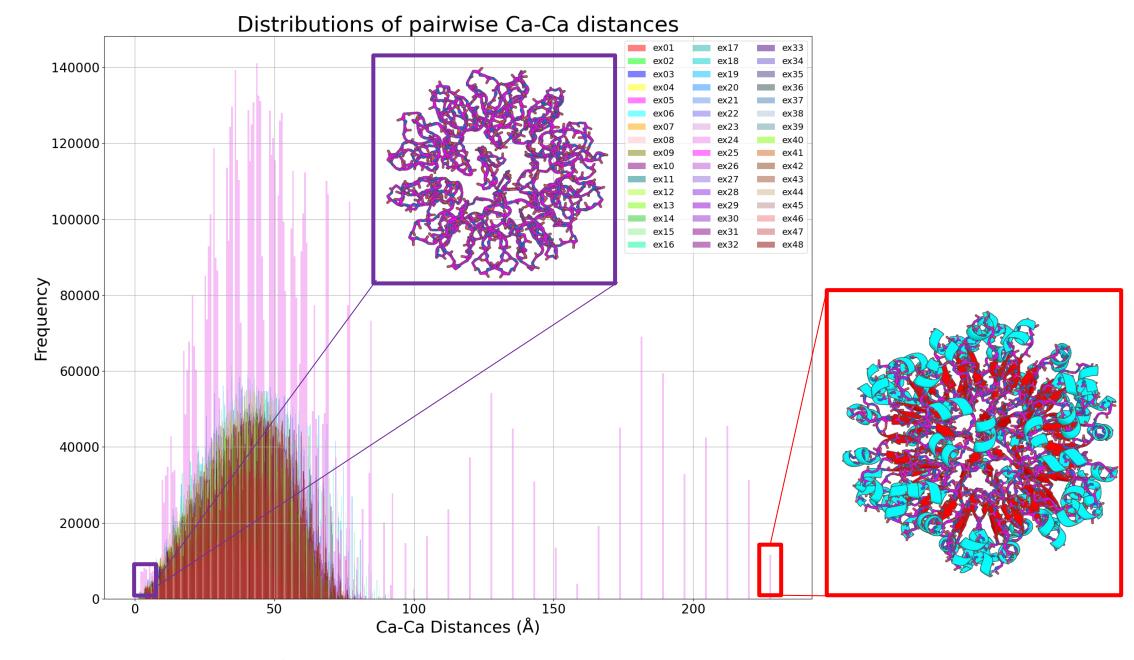
AfCycDesign

AANAALNGCN AANAALNGCN

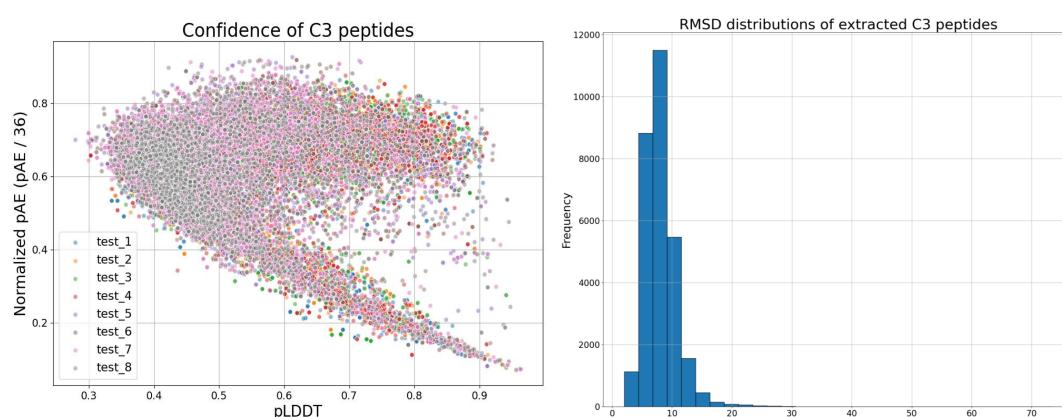
AANAALNGCN.

Results

Sample particles of various packing densities and sizes



Evaluation of extracted cyclized peptides



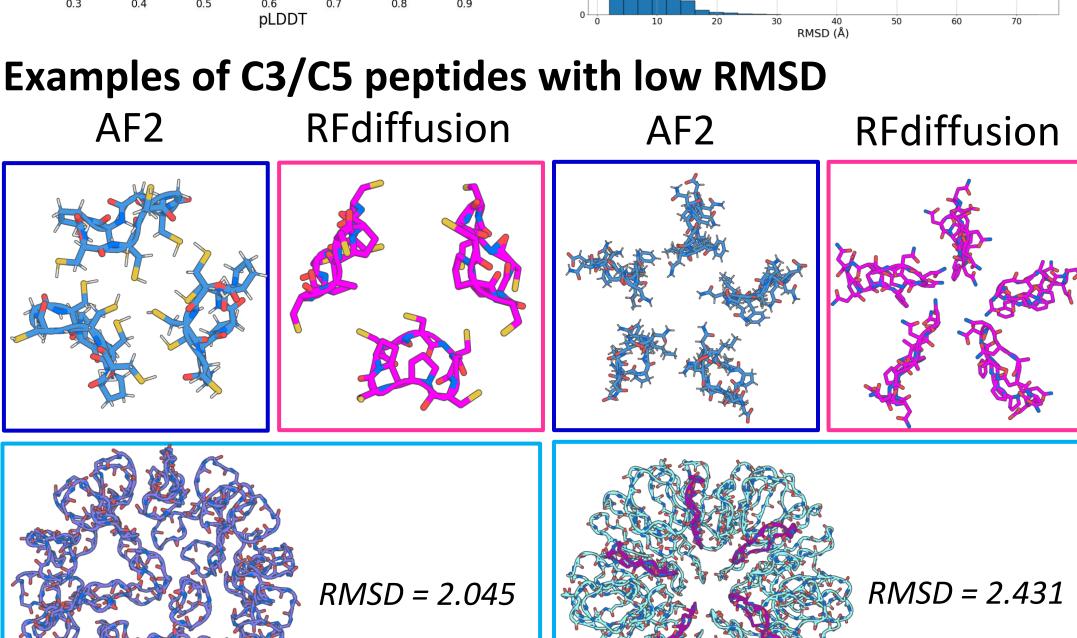


Figure 4. Predictions of C3 and C5 peptides extracted from the icosahedral complex.

Generate new cyclized Design symmetric oligomers via sequences via **RFdiffusion ProteinMPNN** X_t (input) $\widehat{X_0}$ (prediction) AANAAL**Ń**GCN

Hyperparameters

Guiding potential strength/schedule

Chain length

AANAALNGCN Example. Designing symmetric homooligomer sequences with identical side

Symmetric Sequence Design

chains using the "Tied" MPNN model.

Design a symmetric amino acid sequence using ProteinMPNN.

Confident Metrics

- pLDDT > 90
- Normalized pAE < 0.2
- RMSD < 2

Discussion & Progress

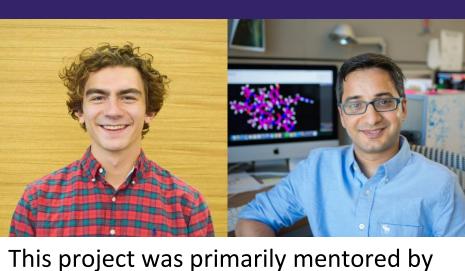
Takeaway

- 1. Icosahedral symmetric oligomers composed of cyclic peptides can be designed with RFdiffusion and validated *in silico* with AfCycDesign⁵.
- 2. Oligomers composed of longer cyclic peptide chains tend to have higher AF2 confidence.

Progress & Future Directions

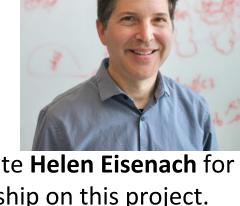
- 1. Characterize promising designs experimentally.
- 2. Generalize the pipeline to other point symmetries (e.g. dihedral and octahedral).
- 3. Change the AF2 settings
 - Use "AF2 initial guess" as in [1].
 - Try running the monomer AF2 weights instead of multimer weights.
- 4. Expand the current pipeline to accommodate scaffolding metal-binding motifs.

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References

[1] Watson, J. et al. (2023). 'De novo design of protein structure and function with Rfdiffusion'. Nature. [2] Daupras, J. et al. (2022). 'Robust deep learningbased protein sequence design using ProteinMPNN'.

[3] Rettie, S. et al. (2023). 'Cyclic peptide structure prediction and design using AlphaFold'. bioRxiv.

Number of diffusion steps **Figure 2.** Design pipeline of cyclized peptide oligomers.