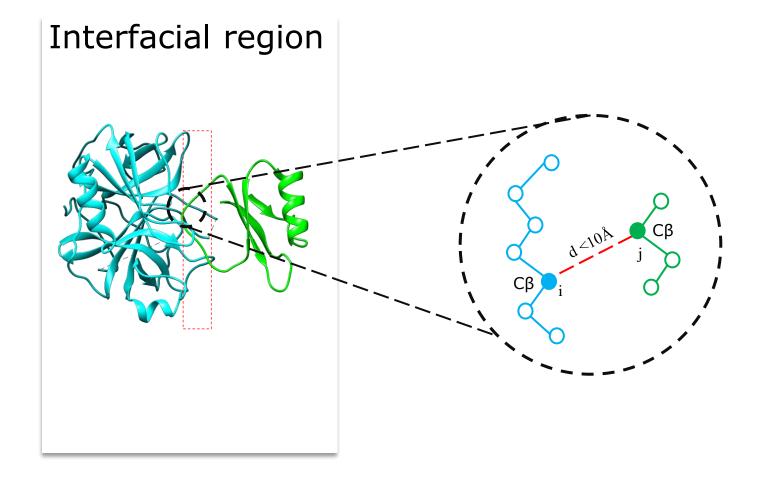
# Estimation of interfacial quality of protein complex models

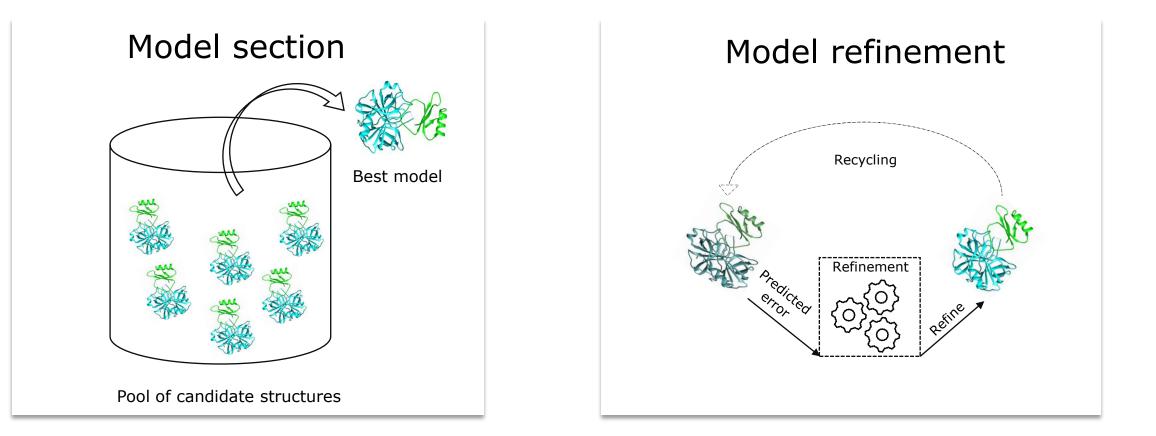
Md Hossain Shuvo Virginia Tech

# Background





# Motivation



Helps in accurately guiding the process of protein complex prediction

# Approach

- Dataset curation
- Feature extraction
- Model training
- Quality estimation

### Dataset

#### Training

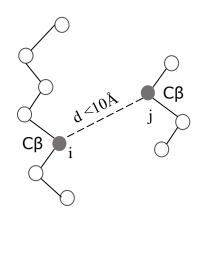
- DOCKGROUND docking decoy set 2
- > 180 complex targets
- ~18000 docking decoys

#### Testing

- DOCKGROUND docking decoy set 1
- > 23 complex targets
- ➤ ~2600 docking decoys

# **Feature extraction**

#### Graph representation G(nodes, edges)



#### Node features

- Amino acids encoding
- Secondary structure and solvent accessibility encoding
- Relative residue positioning
- MSA based features (NEFF)

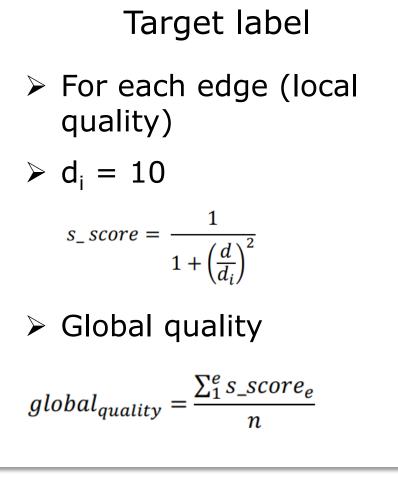
Edge features

- Dihedral angles
  - Phi
  - Psi
- Orientation
  between the
  connecting
  nodes
  - theta
  - Omega
  - phi

# **Quality estimation**

Initial deep learning model

- Graph neural network
- Ideal for learning for graph representation
- Regression problem



## **Evaluation metrices**

#### Ground truth:

- > Observed s-score
- iRMSD (if time permits)
- Pearson correlation between global<sub>quality</sub> and the s-score
- > Spearman correlation between global<sub>quality</sub> and the s-score
- Kendall's Tau correlation between global<sub>quality</sub> and the sscore

# **Competing methods (if time permits)**

- ➢ DOVE
- ➢ GNN-DOVE
- DeepRank

## Milestones

Table1: Tentative timeline for the individual task for the proposed project

Task	Tentative completion date
Data curation	3/11/2022
Feature generation	3/25/2022
Initial network construction	4/1/2022
Network training	4/8/2022
Evaluation	4/13/2022

## Acknowledgement



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