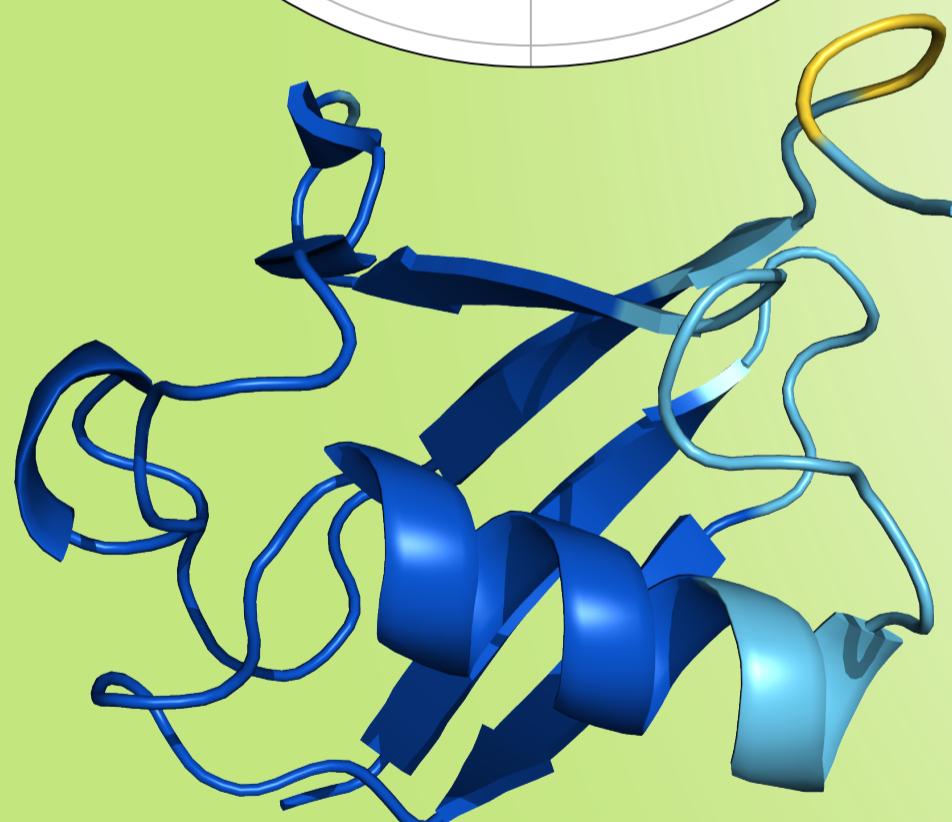
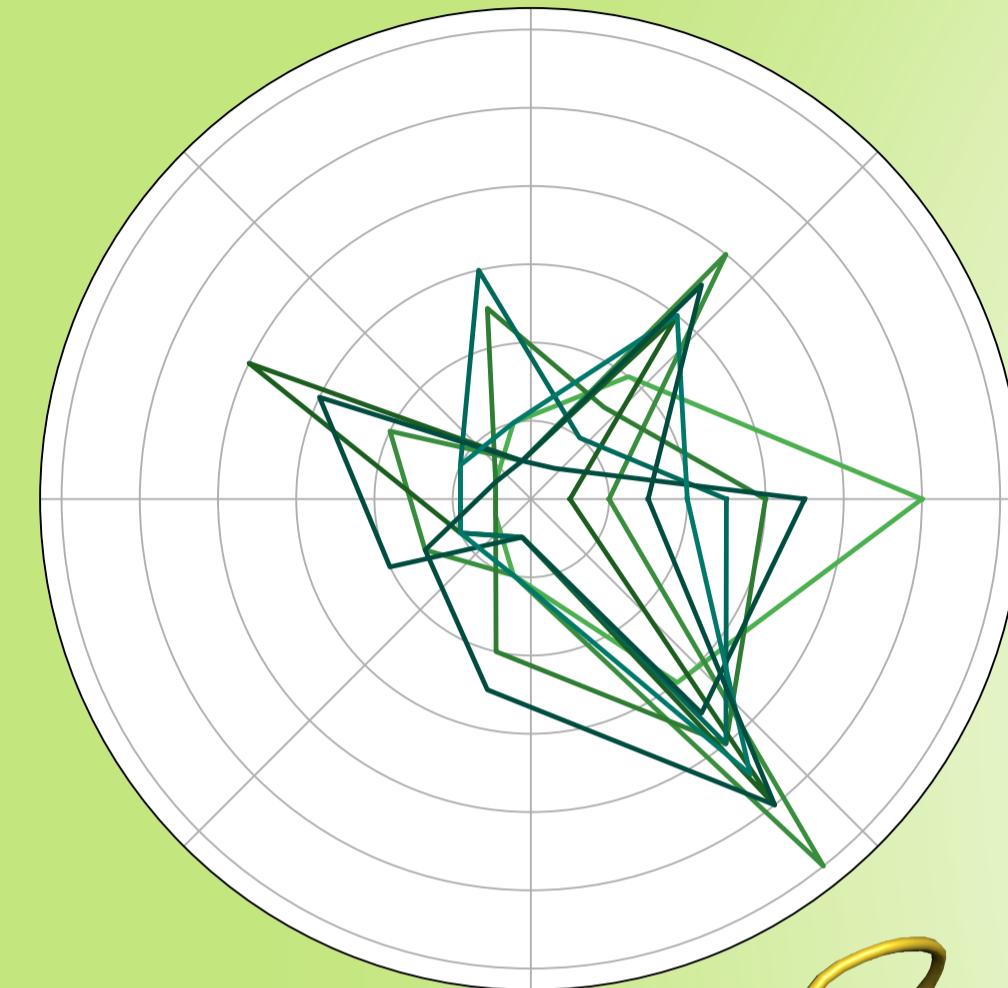
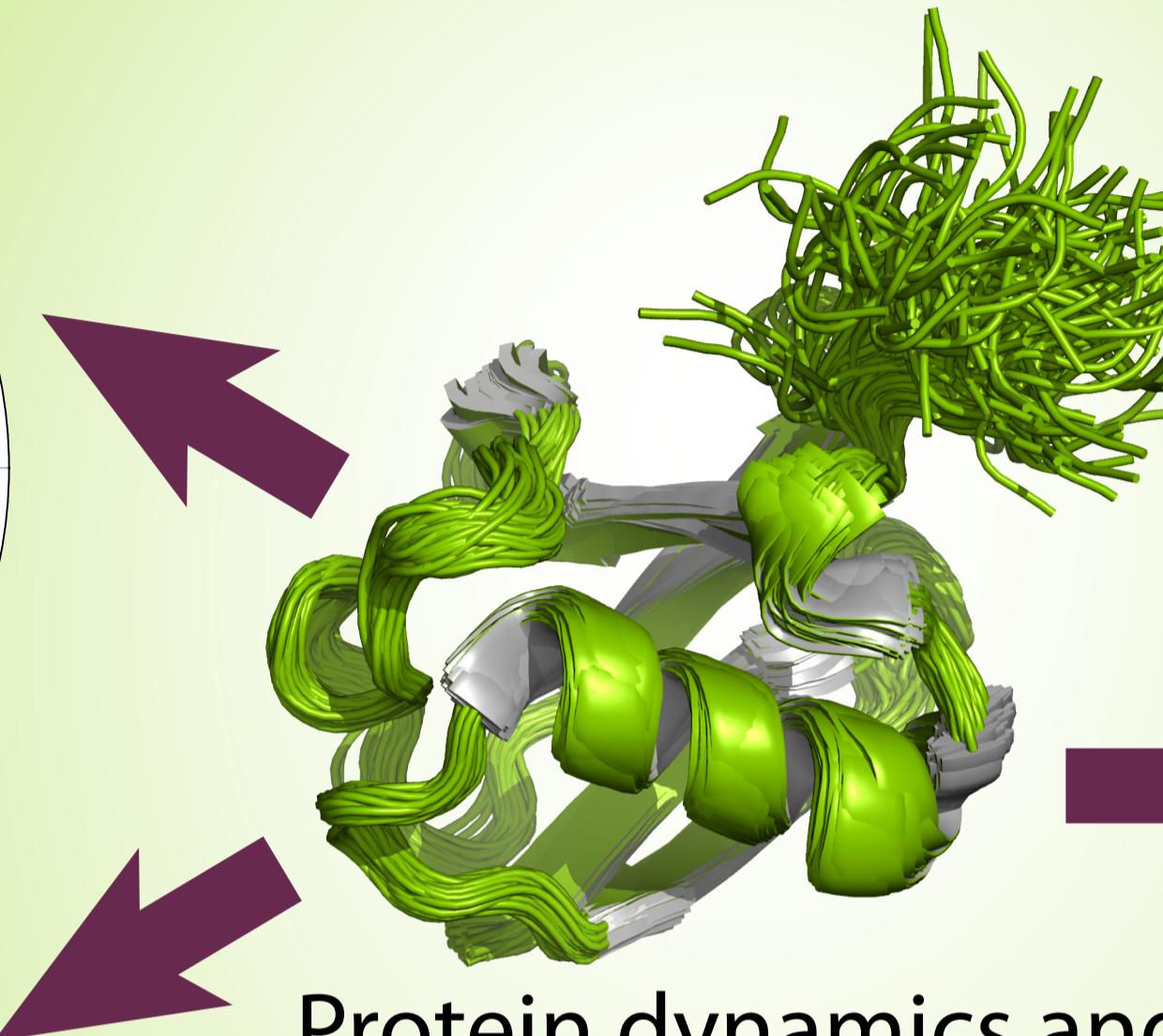


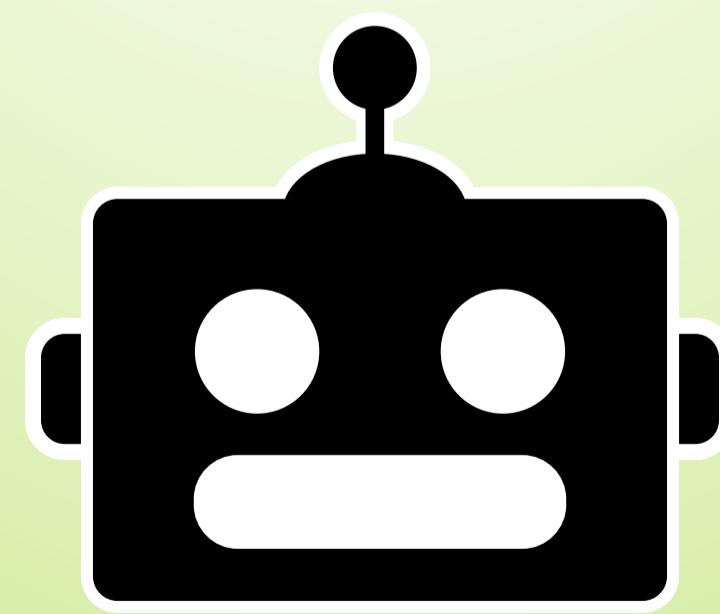
# Statistical interpretation of local conformational states



Investigation of relationship  
between AlphaFold2 models and  
observed and predicted dynamics



Protein dynamics and  
multiple conformations



Predictions as features for models  
on protein ambiguous behaviour

# Predictors deployment

>6V5D|Ubiquitin|Homo sapiens (9606)  
MQIFVKTLTGKTITLEVEPSDTIENVKAKIQQ  
KEGIPPDQQQLIFAGKQLEDGRTLSDynIQK  
ESTLHLVLRLRGG

