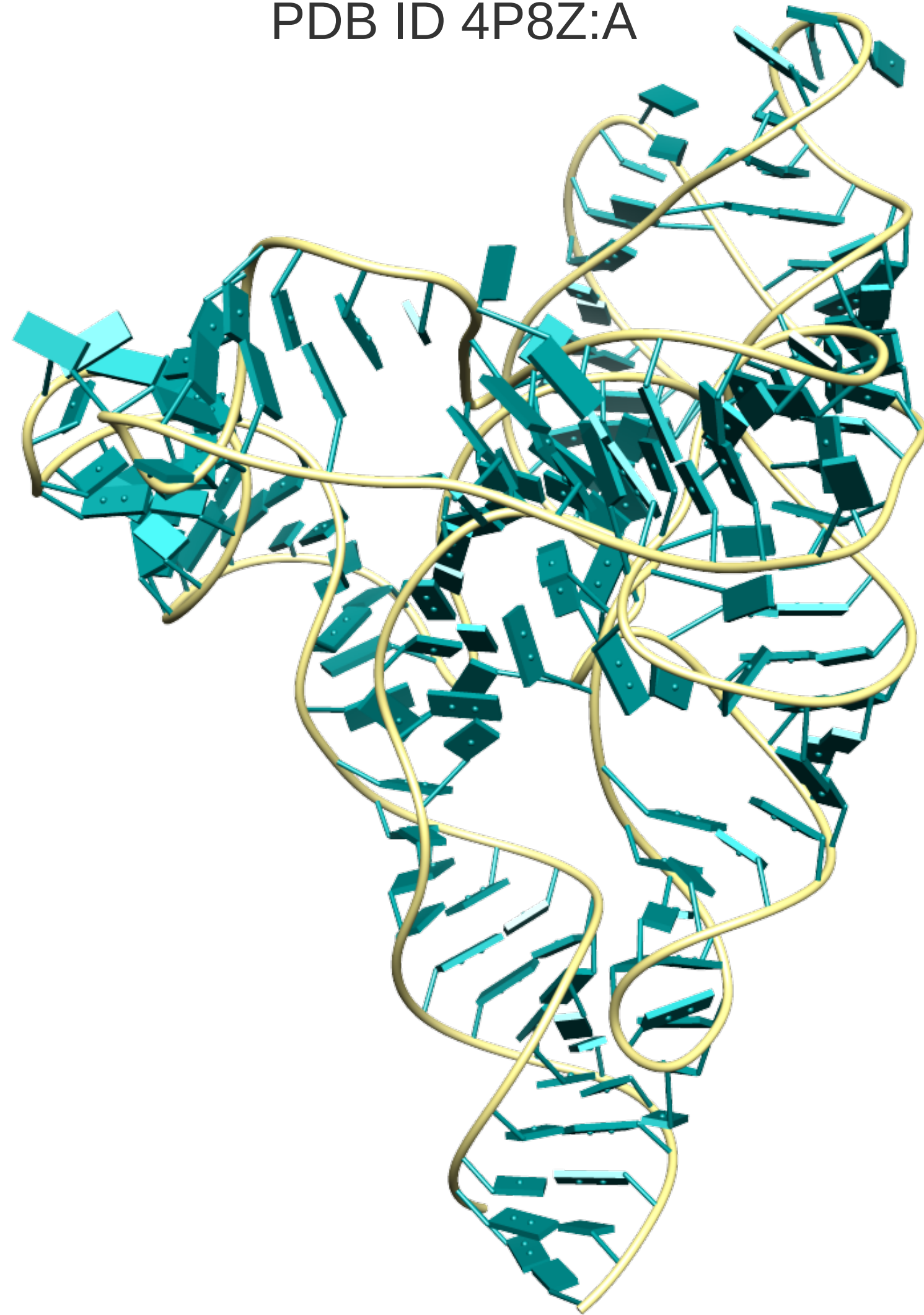
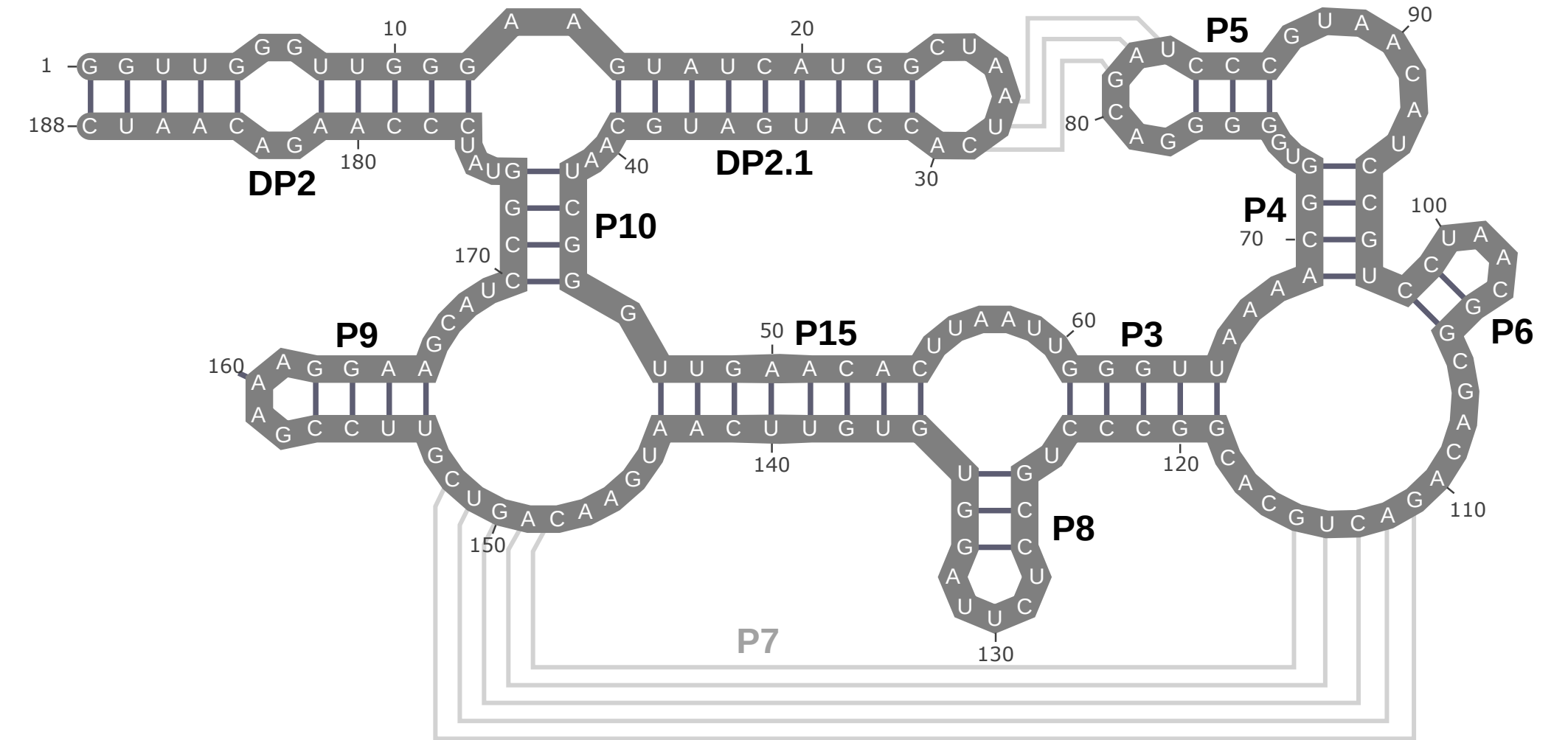


**A** 3.85Å DiLCrz model  
PDB ID 4P8Z:A



**B** Reference 2D structure



**C** Best prediction  
MCC=0.853

Base pairs  
— Correctly predicted  
— Incorrectly predicted  
— Missing

