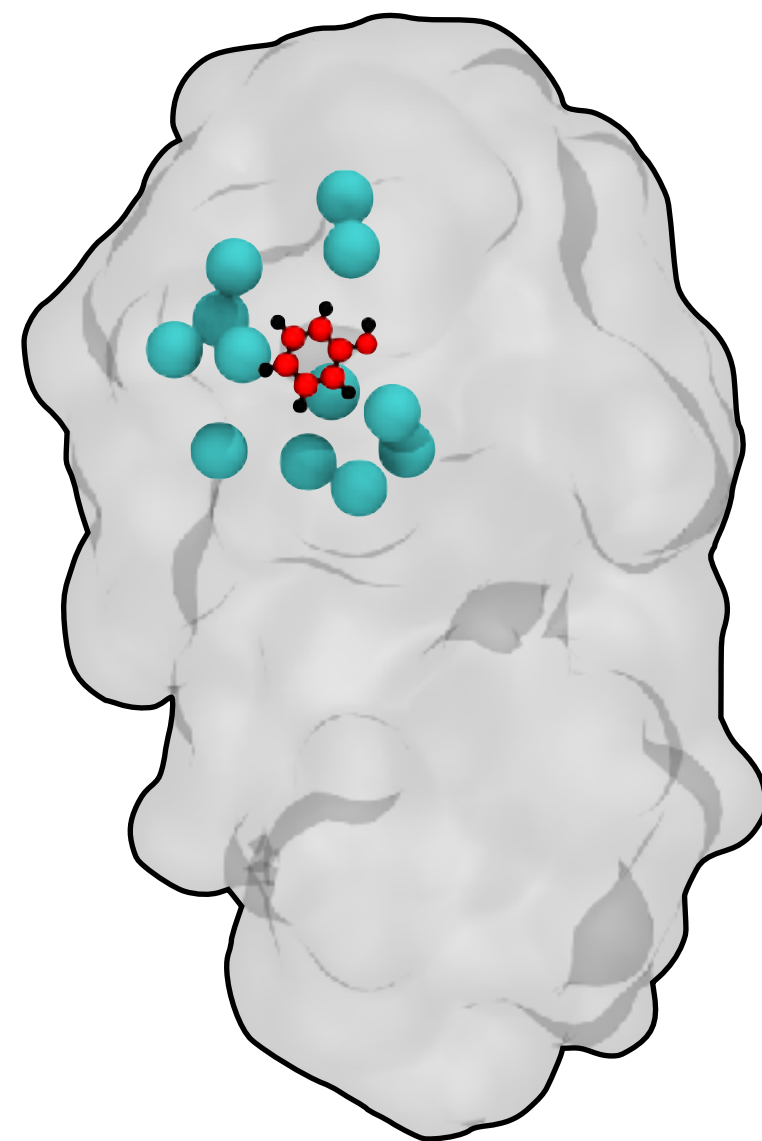
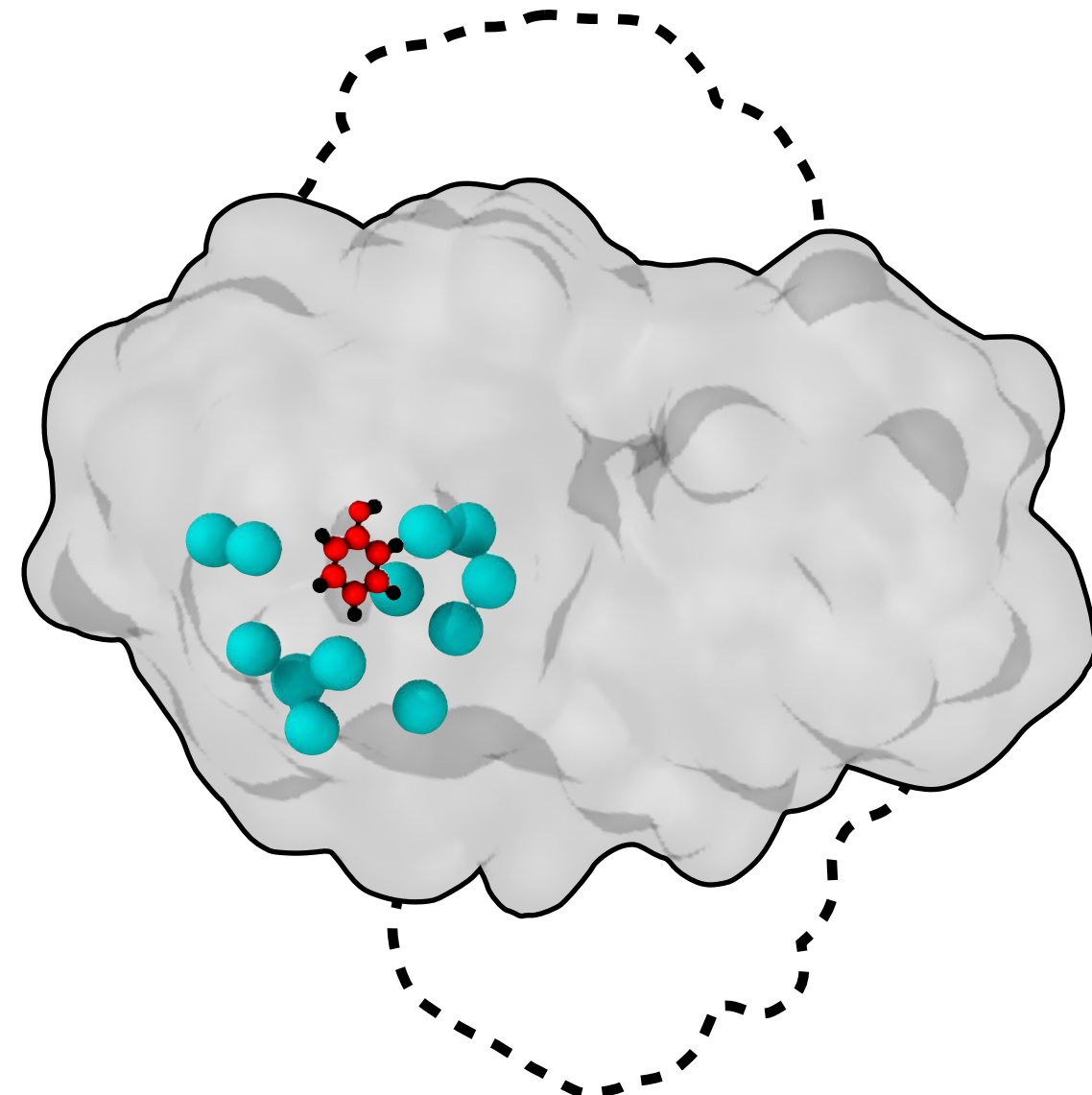
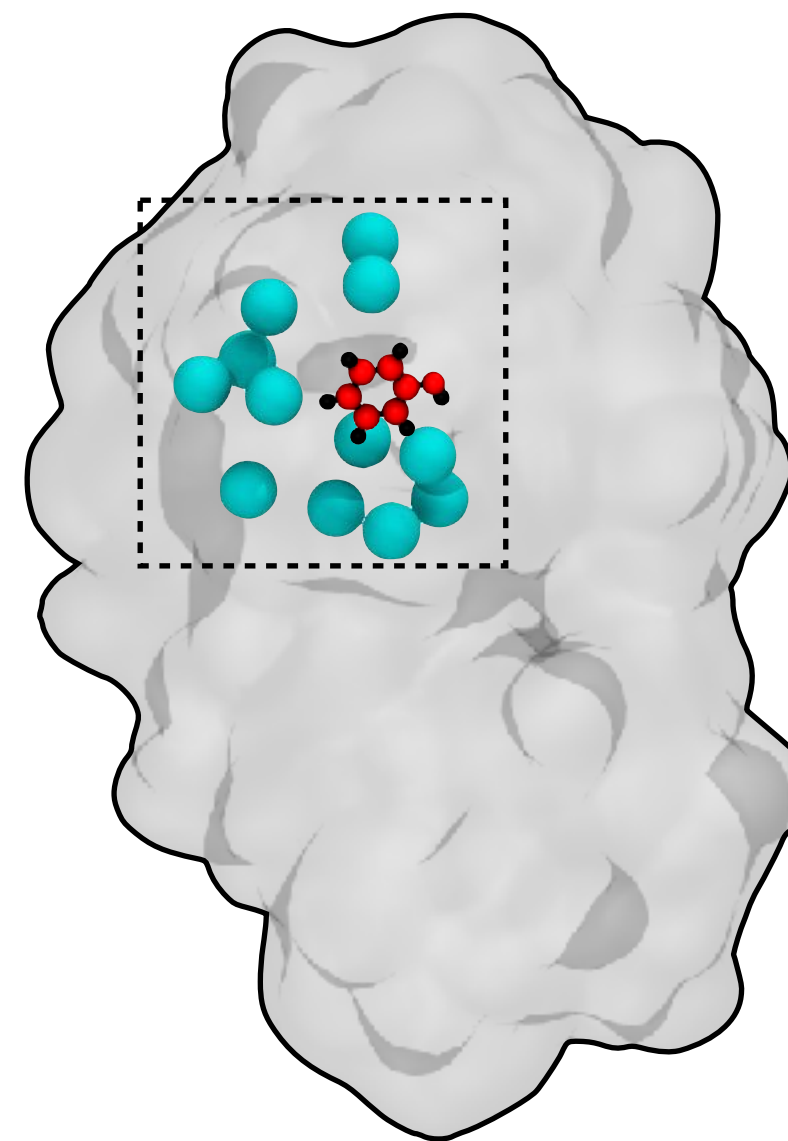
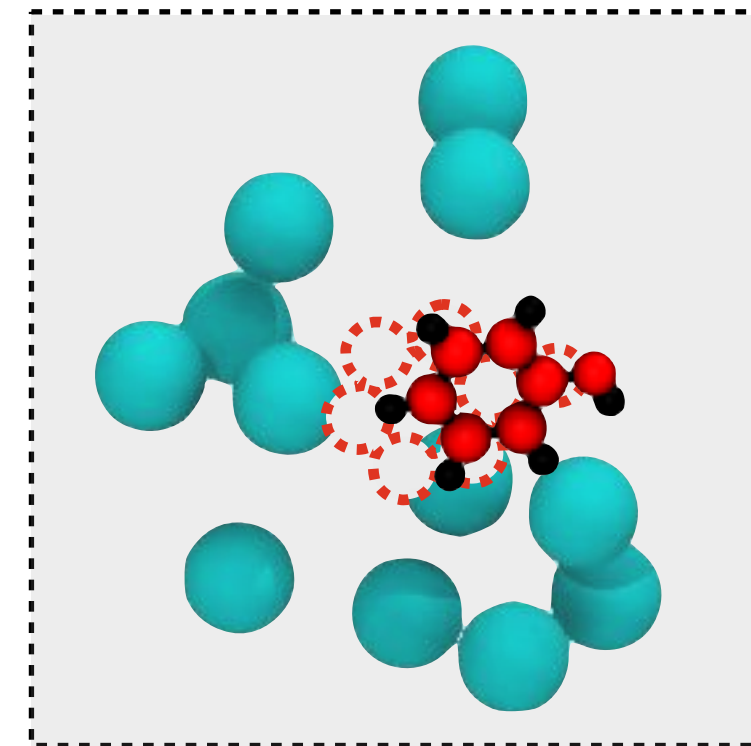


**A****B**

Align to reference  
using **protein**  
**fitting group**

**C**

Measure **ligand**  
RMSD with respect  
to reference

**D**

**Reference Coordinates**

**Calculation of the Instantaneous DBC**