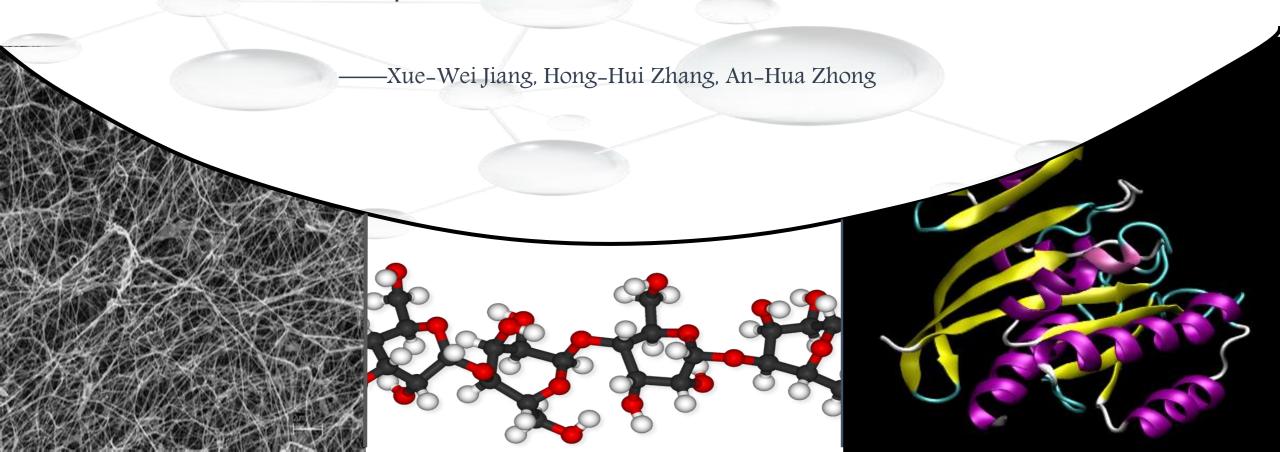
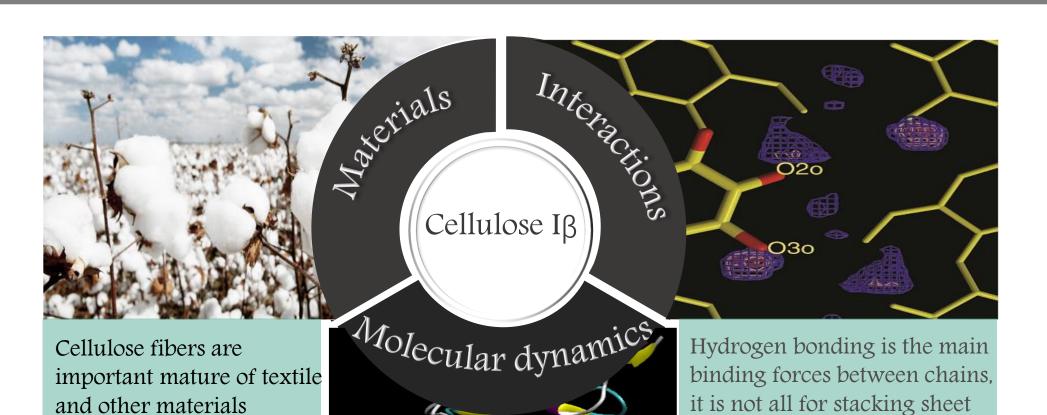


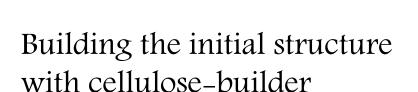
## The Van der Waals and Electrostatic Interaction Studying of

Crystalline Cellulose IB Based on Molecular Dynamics

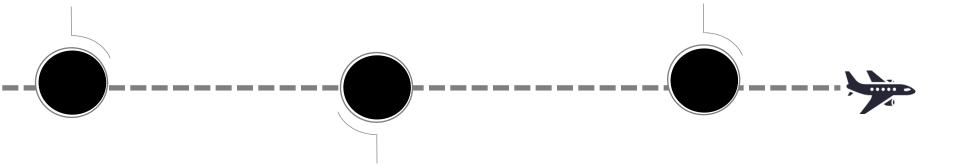




Molecular dynamics approach is an effective computer modeling technology



Calculate the interaction energy by program MMPBSA.py of AmberTools

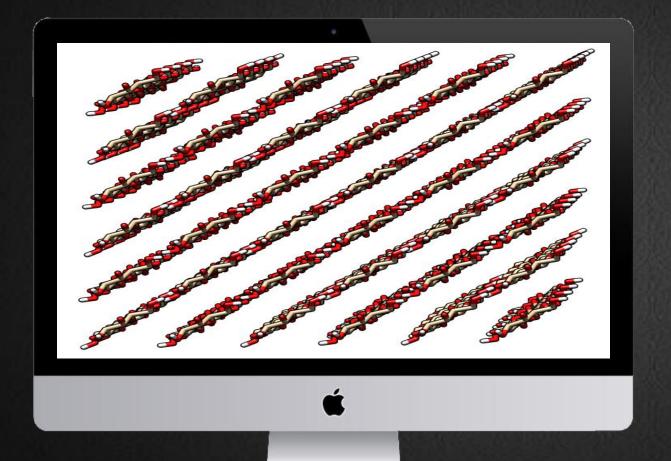


All simulation were performed with Amber software and explicit water



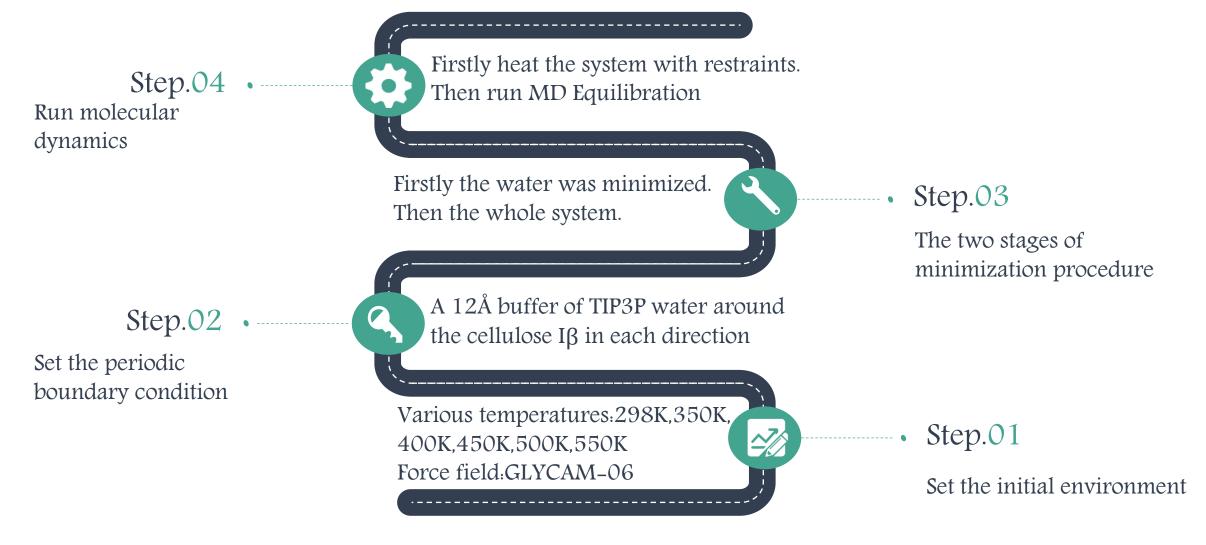
## About the initial modeling

02. Methodology and Processing



- $\Rightarrow$  a = 7.784 Å,b = 8.201 Å,
  - c = 10.38 Å
- $\triangleright$   $\alpha = \beta = 90^{\circ}$  ,  $\gamma = 96.5^{\circ}$
- > c is the chain direction



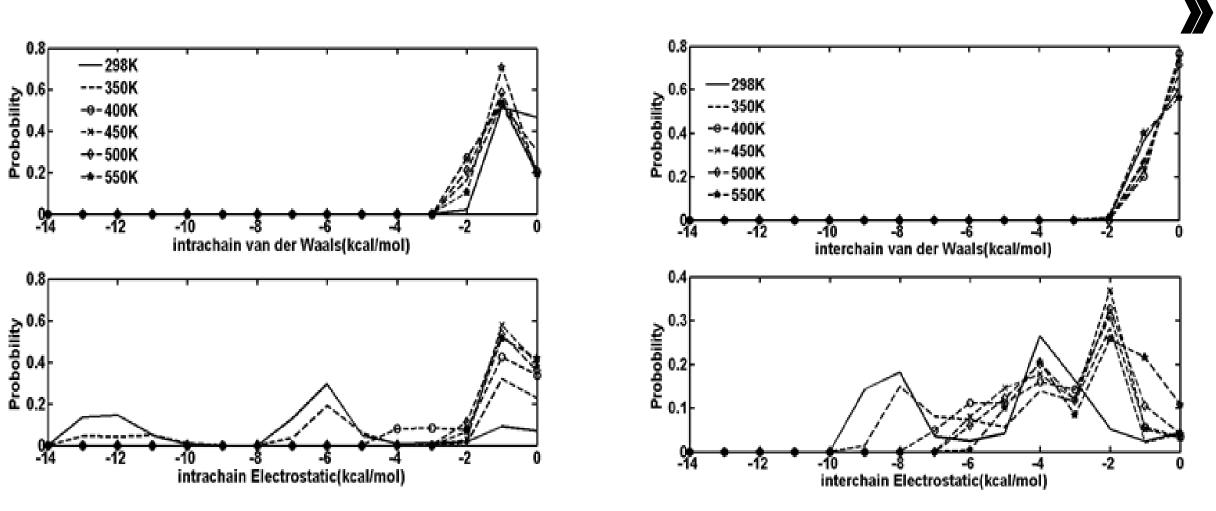


The simulation processing with software Amber



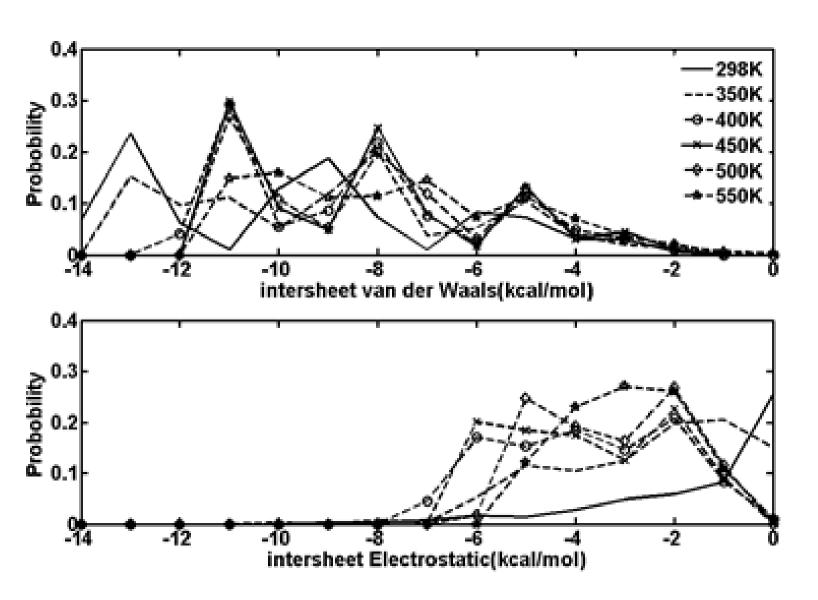
At 298 K, the Van der Waals and electrostatic intermolecular energy per chain are **-43.91 kcal/mol and -26.24** kcal/mol respectively. The Van der Waals energy gradually reduces when the temperature rises, meanwhile the electrostatic energy increases initially and then decreases.



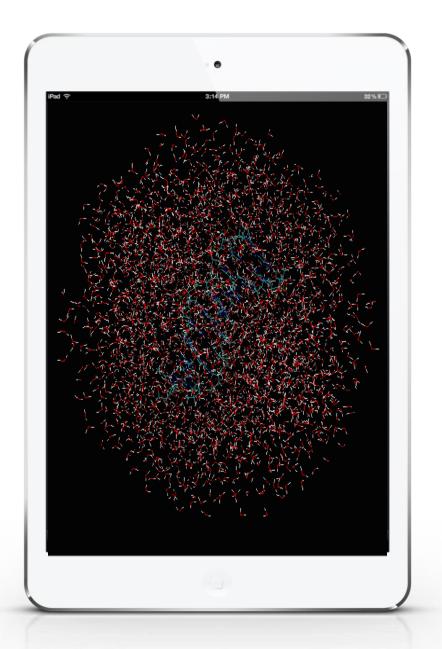


The electrostatic energy of most glucose residues for intrachain lies on 11–14 kcal/mol and 5–8 kcal/mol, while located on 7–10 kcal/mol and 2–5 kcal/mol at 298 K for interchain. The Van der Waals interaction energy in intrachain increases slightly as the temperature rises and there is almost no growth for interchain.





The intersheet Van der Waals interaction decreases slightly with the temperature increasing, The Van der Waals energy of most glucose residues is located on the range of 6–12 kcal/mol at 550 K.



The interactions were discussed only in water, and no comparison to other solvents. In the next step, we will study the Van der Waals and electrostatic interaction in different solvent environments.





THANK YOU.