The first hydration layer around biomolecules is site-specific

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Computational Approaches to RNA Structure and Function

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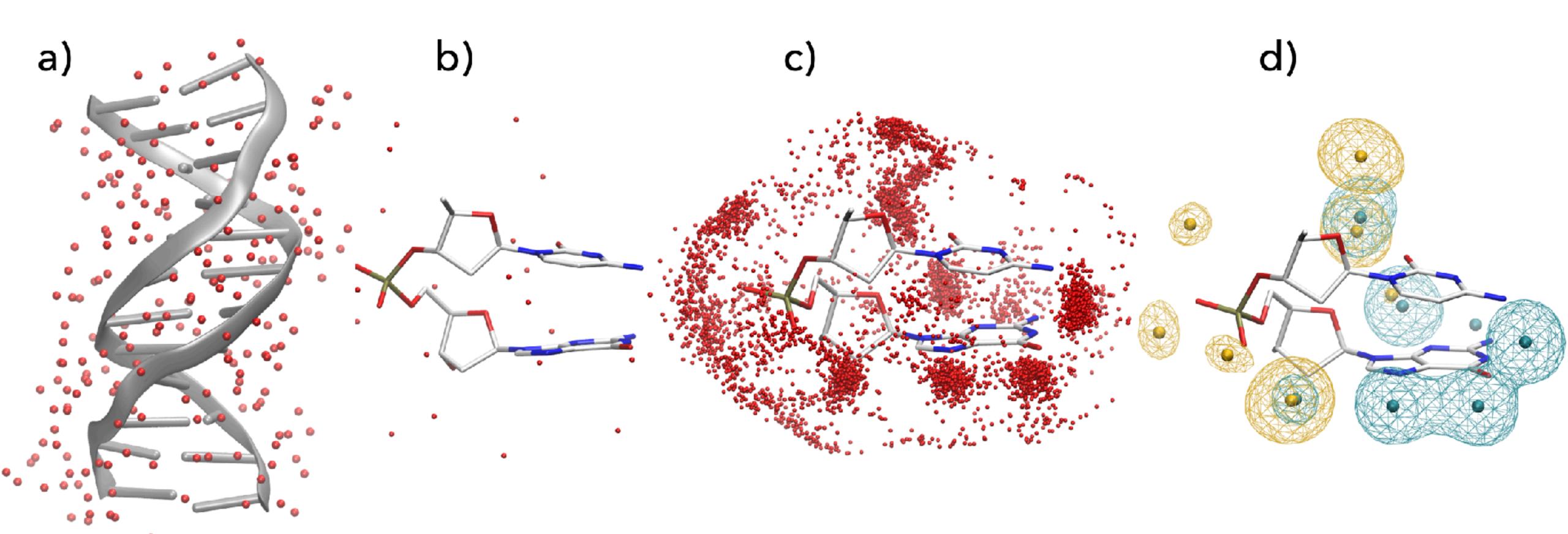


Background

- Proteins and nucleic acids evolved in the aqueous environment.
- The 1st hydration shell has properties different from the bulk water.
 - Biedermannová L. & Schneider B.: Hydration of proteins and nucleic acids: Advances in experiment and theory. A review. Biochimica et Biophysica Acta General Subjects 1860: 1821-1835 (2016).
- Water dynamics in the hydration shell is reduced compared to the bulk.
 - Bohdan Schneider, Jean-Christophe Gelly, Alexandre G. de Brevern & Jiří Černý: Local dynamics of proteins and DNA evaluated from crystallographic B factors. Acta Cryst. D70: 2413–2419 (2014).
- The first hydration shell consists of mostly ordered (localized) water molecules.
- Proteins:
 - Biedermannová L. & Schneider B.: Structure of the ordered hydration of amino acids in proteins: analysis of crystal structures. Acta Cryst D71: 2192-2202 (2015).
- DNA:
 - Lada Biedermannová, Jiří Černý, Michal Malý, Michaela Nekardová & Bohdan Schneider: Knowledge-based prediction of DNA hydration using hydrated dinucleotides as building blocks. Acta Cryst. D78: 1032–1045 (2022).
- The release of the ordered hydration water is entropically favorable, enthalpically neutral.



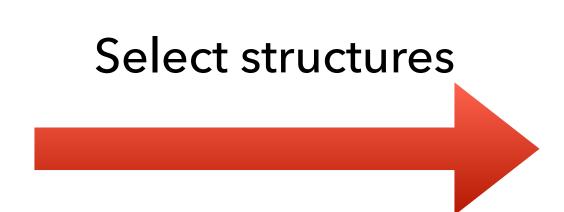
The Protocol

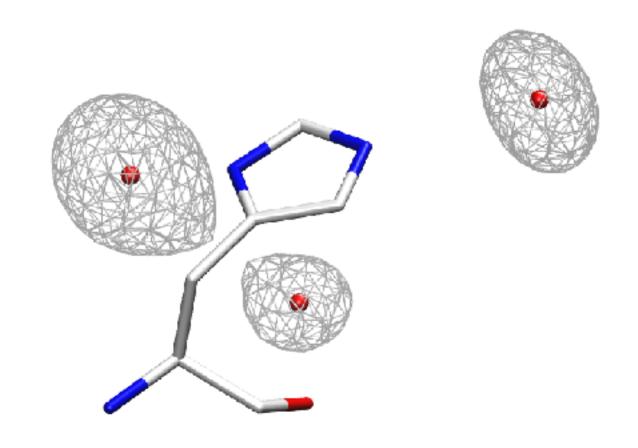






Database search for suitable biomolecular X-ray structures





Fourier Averaging

Hydration sites identified as maxima in water density map

Large number of well-resolved, hydrated X-ray structures Extract hydrated fragments Relevant water molecules

added to a reference fragment

The Key Point of Our Hydration Analysis

- You rely on sensibly selected fragment
- When the fragment has conformational freedom, you must have well clustered conformers
 - rotamers for amino acids
 - NtC dinucleotide classes for DNA and RNA



To Be Finished

- Interpretation of RNA hydration patters
- RNA hydration to watna



The website

• Černý J., Schneider B. & Biedermannová L.: WatAA: Atlas of Protein Hydration. Exploring synergies between data mining and ab initio calculations. Phys. Chem. Chem. Phys. 19, 17094 (2017).



