

Observing solvation free energy changes in biological processes with THz spectroscopy

Simone Pezzotti

Ruhr University Bochum

Solvation in biology: importance & challenges

Protein folding



D. M. Huang et al. **PNAS**, 97, 8324 (2000).

Hydration entropy & enthalpy

Liquid-liquid phase separation



S. F. Banani et al. Nat. Rev. Mol. Cell Biol. 18, 285 (2017).

Solvation in biology: importance & challenges

Protein folding



V. C. Nibali et al. JPCL, 11, 4809 (2020)

Hydration entropy & enthalpy

Liquid-liquid phase separation



Challenges Map local solvation contributions Improve time resolution

N. B. Rego et al. PNAS, 118, e2018234118 (2021)

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Protein folding



Hydration entropy & enthalpy

Liquid-liquid phase separation





N. B. Rego et al. PNAS, 118, e2018234118 (2021)

THz spectral range is ideal to look at hydration



Advantages of THz:

- ✓ Hydration shell motions
- ✓ Distinguish water/solute

THz spectral range is ideal to look at hydration



F. Bohm et al. Angewandte 56, 9981 (2017)





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Quantities
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Can be recorded as a fiction of time down to ps resolution

1. Create cavity









$$\Delta G_{solv} = \Delta G_{cavity} + \Delta G_{bound}$$



THz fingerprints of the 2-steps: 1. cavity-wrap



Spectral dissection from theoretical spectroscopy (DFT-MD)

V. C. Nibali et al., **JPCL**, 11, 4809 (2020). <u>S. Pezzotti</u> et al. **Angewandte**, e202203893 (2022)

THz fingerprints of the 2-steps: 1. cavity-wrap



Cavity-wrap = Hydrophobic



Collective HB-stretching

V. C. Nibali et al., **JPCL**, 11, 4809 (2020). <u>S. Pezzotti</u> et al. **Angewandte**, e202203893 (2022)

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Bound = Hydrophilic



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Hindered rotations

V. C. Nibali et al., **JPCL**, 11, 4809 (2020). <u>S. Pezzotti</u> et al. **Angewandte**, e202203893 (2022)





V. C. Nibali et al., **JPCL**, 11, 4809 (2020). <u>S. Pezzotti</u> et al. **Angewandte**, e202203893 (2022)







$$\Delta G_{cavity} = \Delta \alpha_{wrap} \Delta \overline{H}_{wrap} - T \Delta \alpha_{wrap} \Delta \overline{S}_{wrap}$$
$$\Delta \overline{G}_{bound} = \Delta \alpha_{bound} \Delta \overline{H}_{bound} - T \Delta \alpha_{bound} \Delta \overline{S}_{bound}$$



Measured

Wrap/bound THz fingerprints



✓ Accurate solvation free energies



✓ Accurate solvation free energies



1.0

✓ Accurate solvation free energies



✓ Accurate solvation free energies

✓ Molecular interpretation



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S. Pezzotti et al. Angewandte, e202203893 (2022)

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S. Pezzotti et al. Angewandte, e202203893 (2022)

✓ Accurate solvation free energies

✓ Molecular interpretation

✓ Watch free energy changes in real-time



S. Pezzotti et al. JPCL, 14, 1556 (2023)





- **Biological relevance**: membran-less compartmentalization
- **Medical relevance**: pathological protein aggregates



- **Biological relevance**: membran-less compartmentalization
- Medical relevance: pathological protein aggregates



$$\delta G_{LLPS} = \Delta G_{p-p} + \Delta G_{cavity} + \Delta G_{bound}$$









J. Ahlers et al. **Biophys. J.** 120, 1266 (2021) <u>Pezzotti</u> et al. **JPCL**, 14, 1556 (2023)





J. Ahlers et al. **Biophys. J.** 120, 1266 (2021) <u>Pezzotti</u> et al. **JPCL**, 14, 1556 (2023)



J. Ahlers et al. **Biophys. J.** 120, 1266 (2021) <u>Pezzotti</u> et al. **JPCL**, 14, 1556 (2023)





- Wrap (Hydrophobic) water is released
- Bound (Hydrophilic) water is retained





• We can determine if LLPS takes place



- We can determine if LLPS takes place
- The individual terms are huge, but compensate.





Tuning the hydrophilic/hydrophobic balance









Sebastiani, <u>Pezzotti</u>, et. al. **PNAS**, 52, 32954 (2020)

Different from any other water phase



Confined water release = entropic driving force

Sebastiani, Pezzotti, et. al. PNAS, 52, 32954 (2020)





Regulate guest encapsulation



Sebastiani, <u>Pezzotti</u>, et. al. **PNAS**, 52, 32954 (2020)

Conclusions on THz calorimetry

- We can directly correlate solvation structure-spectroscopy-thermodynamics
- We can dissect local hydrophobic & hydrophilic hydration contributions



Conclusions on THz calorimetry

- We can directly correlate solvation structure-spectroscopy-thermodynamics
- We can dissect local hydrophobic & hydrophilic hydration contributions



• We can reveal solvation driving forces to catalytic and biological processes in real-time





F. Sebastiani, et al., **PNAS**, 2020, 117, 32954

Acknowledgments



RUB

Collaborations

J. Tatzelt (RELOLV) K. F. Winklhofer (RELOLV) I. Sagi (Israel) M. Heyden (U.S.) V. Conti Nibaly (Italy) F. Sebastiani (Italy) M.-P. Gaigeot (France) Songi Han (U.S.) S. Shell (U. S.)





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Thank you for your attention





