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#### Winter College on Micro and Nano Photonics for Life Sciences

11 - 22 February 2008

THz spectroscopy of proteins

Martina Havenith Ruhr University Bochum Bochum, Germany



#### The THz dance of water with biomolecules M. Havenith Physical Chemistry, Ruhr-Universität Bochum, Germany

#### **Funding:** Human Frontier Science Award 2004

### Physical Chemistry: How it started



Robert Wilhelm Bunsen chemist (1811-1899) Bunsen: 15. 11. 1859:

"At the moment Kirchhoff and I are busy with a work which keeps us awake all night. Kirchhoff has made an unexpected wonderful discovery, by finding the reason for the dark lines in the spectrum of the sun."



Gustav Kirchhoff physicist, mathematician (1824-1887)

Chemische Analyse durch Spectralbeobachtungen (1860) Abhandlungen über Emission und Absorption (1859-1862)



### Why did it take so long ?



Sir John Frederick Herschel (1792 - 1871)

John Frederick Herschel Physiker, Mathematiker (1827)

"Their color would be a good means to detect even small quantities of dyed materials. But since this is chemistry one doesn't talk about it !"





"There is no science which is lacking in soul, with more inability to think, a greater lack of true insight and understanding, more shortsighted and weakness under the cover of knowledge and erudition as mathematics."



Baron Justus von Liebig (1803 - 1873)

# Taking a closer look into the world of biomolecules: Spectral Fingerprints

Skeleton motions Breathing modes Large amplitude vibrations





MIR

avelength

energy

**Determination of structure** 

#### The THz gap between electronics and photonics



Figure 2. Illustration of the real THz gap-the decline in power as electronics goes to higher frequencies, and photonics goes to lower frequencies. Electronic data courtesy of Tom Crowe, Virginia Diodes Inc.

#### **Entering the THz age**

• Electronic: Acceleration of charges limited by speed of charge accelaration THz corresponds to fsec

 Photonic: transition between conduction band and valence band limited by possible band gap in semiconductors



Laser transition

- The new radiation sources:
  - **p-Ge laser THz time domain spectrometer (TDS)** QCL quantum cascade laser (imaging ) photomixer Free electron laser

## THz spectroscopy a new spectral window for applications in chemistry, medicine and material science

#### T-Ray: Next frontier in Science and Technology

Terahertz wave (or T-ray), which is electromagnetic radiation in a frequency interval from 0.1 to 10 THz, lies a frequency range with rich science but unexplored technology.





#### Publications from 1985-2005

#### The new sources: **Tunable THz Germanium Laser - Emission** - a high power laser in the THz region-



#### A microscopic view on biomolecules



Do we have a molecular description of macroscopic phenomena ? *"How do physical forces and interactions govern biological function, from the molecular to the cellular scale ?"* 

Can we predict the function of the protein based upon merely the structure ?

Additional relevant parameters ? Dynamics? Role of water ?

Can we find adequate models for theoretical description of solvation?

#### Solvation of proteins:

Expulsion of solvation water the hydrophobic core cause folding barrier Dynamical and structural fluctuations of water molecules contribute to the flexibility of the polypeptide chain : e.g. heme binding site Slaving? Do solvent fluctuations dominate protein function ? How does solvation water differ from bulk water ? What is the size of the hydration layer ?



#### THz absorption of water yields information on network oscillations – solvation dynamics

Beers lawResult:f = 2.7 THz $I(\omega) = I_0 \cdot e^{-\alpha(\omega)I} + \text{ const.}$  $\alpha \approx 400 \text{ cm}^{-1}$ ! $g = 90 \text{ cm}^{-1}$ 

#### Strong temperature dependence of THz absorption

Ronne, Thrane, Astrand, Wallqvist, Mikkelsen, Keiding, J. Chem. Phys. 107, 5319 (1997)

Water absorption at 1.5 THz for 100 µm water layer:





370 K :  $I/I_0 = 4 * 10^{-1} = 40 \%$ 



Molecular dynamic simulationen (MOIL) X. Xu, D. Leitner, University of Reno water motions at 3 THz (sub-psec dynamic)





Libration (675 cm-1): rupture and breaking of H-bonds



connectivity (200 cm-1): Longitudial motion of the H-bond along the axis

### Anhydrobiosis "life without water"



- Anhydrobiotes can survive complete dehydration
- For plants we find a correlation with the presence of larger amounts of solubale disaccharides, especially trehalose.
- Although this is used in industrial production processes, the molecular mechanism is remains still unclear, so far.

Well known: "Bioprotection" of Liposomes, proteins and red blood cells by sugar Competing molecular description:

- **Water exclusion:** •
  - H-bridge of the sugar to the protein
  - Crowe, Science 223, 701 (1984)



Crowe et al.



Bio-molecule **disaccharide** 

water

crystallisation (Dihydrat Th) **Formation of water channels** Immobilization of water D. Kilburn et al., Nature mater. 5, 632 (2006)





THZ spectroscopy of solvated lactose: Observed: "Bioprotection" of proteins by sugars Hydrogen bonding of sugars to proteins or: Do carbohydrates slow down the dynamics ? How long ranged is the influence of lactose on a psec scale? (Crowe, Science <u>223</u>, 701 (1984), Fayer *et al.* JACS <u>127</u>, 14279)

solvation water

<sup>7</sup>bulk



exchange between bulk and solvent: 10-100 psec

#### *THz spectroscopy* **Tunable THz p-Germanium Laser - Emission** - a high power (2 W) laser in the THz region-



#### THz absorption spectroscopy of solutes Set-up of a p-Ge spectrometer



D. Chamberlain und E. Haller, Rev. Sci. Instrum., 76, 2005

### The THz Spectrum of solvated Lactose

Lactose (gas phase) is nearly transparent in the THz spectral range

#### simple model:

THz absorption is decreased by water displacement









R.A. Jockusch, R.T. Kroemer, F.O. Talbot, L.C. Snoek, P. Çarçabal, J.P. Simons, **M. Havenith,** J.M. Bakker, I. Compagnon, G. Meijer, G. von Helden **J. Am. Chem. Soc. 126, 5709 (2004)** 

#### **Concentration dependence of the THz** spectrum of solvated lactose Soluble (Trehalose) С Hydration layer **Free Water** Lactose concentration **Assumptions:** Excluded volume model: ⇒linear decrease of absorption **Onset of** THz $\alpha_{total} = (1 - V_{solute}/V) \alpha_{bulk}$ nonlinearity Absorption • Each lactose contributes with its is determined α<sub>bulk</sub> THz absorption by the size of the 400 cm-⇒Linear increase + V<sub>solute</sub>/V α<sub>solute</sub> hydration layer Hydration water contributes to total absorption $\Rightarrow$ Linear contribution + V<sub>sw</sub>/V $\alpha_{sw}$ overlap of hydration shells ⇒deviation from linearity

Lactose

### THz spectroscopy of solvated lactose First experimental Results



 $\alpha_{sw}$ : absorption coefficient of the solvation water

 U. Heugen, G. Schwaab, E. Bründermann, M. Heyden, X. Yu, D.M. Leitner, M. Havenith Solute induced retardation of water dynamics: Hydration water probed directly by THz spectroscopy, PNAS 103, 12301 (2006)

# Quantitative description of solvation dynamicsby correlation function

Probability that an H-bond exists at a time t, if it was intact at t=0 C(t)=<h(0)h(t) >/<h> c(t)= exp(-t/ т)





THz spectroscopy as a new tool to probe the dynamical hydration layer (global probe) 123 water per one sugar Determination of the (sub-psec) solvation radius





 $R_{solv}$ = (5.13±0.24) Å Corresponds to 123 H<sub>2</sub>O (much more than X-ray)

Significant retardation up to 6 Å (112 H<sub>2</sub>O) Particullary sluggish for 46 water (4 Å )

#### THz difference absorption spectrometer for precise measurements of absorption coefficients





Comparison between different sugars: scaling the change of THz absorption with the number of H-bonds ((n<sub>c-w</sub>-HB))



### THz spectroscopy as a tool to determine the size and the "nature" of the hydration shell in the sub-psec regime

Fit to a three component model:

 $\alpha_{total} =$ 

$$V_{sw}/V \alpha_{sw} + V_{solute}/V \alpha_{solute}$$
  
+ (1-V<sub>sw</sub>/V-V<sub>solute</sub>/V)  $\alpha_{bulk}$ 

 $V_{sw}/V$ : relative volume of the solvation water  $\alpha_{sw}$ : absorption coefficient of the solvation water  $V_{solute}/V$ : relative volume of the solute  $\alpha_{solute}$ : absorption coefficient of the solute

# Comparison of the dynamical hydration shell for different carbohydrates





Trehalose R<sub>sol</sub>= (6.5±0.9) Å ~ 190 H<sub>2</sub>O



Lactose R<sub>sol</sub>= (5.7±0.4) Á ~ 150 H<sub>2</sub>O

Glucose R<sub>sol</sub>= (3.7±0.9) Å ~ 50 H<sub>2</sub>O

#### THz absorption of bulk and hydration water Hydration water shows increased THz absorption



Conclusions by comparison to MD:

Network vibrations of water molecules near lactose oscillate more in phase than bulk water

Lactose and hydration water molecules oscillate coherently

Coherent ocsillation yields increased THz absorption

U. Heugen, G. Schwaab, E. Bründermann, M. Heyden, X. Yu, D.M. Leitner, M. Havenith, PNAS **103**, 12301 (2006)

#### Mol. Dyn. simulation (3nm x 3nm x 3nm box) M. Heyden, U. Heugen, G. Nienhues, E. Bründermann, D. Leitner, M. Havenith, JACS under review



#### Molecular dynamic simulation (MOIL) X. Xu, D. Leitner, University of Reno Motions at 3 THz (sub-psec Dynamik)



Lactose and Hydration water



collective motions of water molecules in the network

### Individual versus collective oscillations



#### **Bulk water: Uncorrelated, incoherent motion**

#### Individual versus collective motions



#### Hydration water: Coherent, in phase motion

### How about proteins? What absorbs more proteins of the water ? Do we see "biological water" ?



# Taking a closer look on protein and water dynamics:

D.M. Leitner, M. Havenith M. Gruebele Intl. Reviews in Physical Chemistry 25, 553 (2006)



molecular dynamics simulation (MOIL) and figures by: X. Xu, D. Leitner

#### **Previous study on THz spectroscopy of proteins**

Probing the collective vibrational dynamics of a protein in liquid water by terahertz absorption spectroscopy

JING XU,<sup>1</sup> KEVIN W. PLAXCO,<sup>2</sup> AND S. JAMES ALLEN<sup>1</sup> Departments of <sup>1</sup>Physics and <sup>2</sup>Chemistry and Biochemistry, University of California, Santa Barbara, California 93106, USA (RECEIVED January 2, 2006; FINAL REVISION February 13, 2006; ACCEPTED February 16, 2006)



THz absorption of BSA solutions at 1.56 THZ

#### Highest concentration in our experiment

#### Protein science 15, 1175 (2006)

Conclusion: Linear decrease with protein Concentrations Proteins absorb less than water

Could deduce the spectrum of the THz protein : linear increasing with frequency

Possible coupling of hydration dynamics to protein dynamics (Whitmore, Wolpert, Markelz et al. Biophys. J. 85, 1269 (2003))

### THz absorption of solvated lamda-repressor Experimental results:

#### non monotonic concentration dependence !

![](_page_36_Figure_2.jpeg)

M. Gruebele, D. Leitner, and M. Havenith, PNAS 104, 20749 (2007)

#### The THz dance of the water with the protein

![](_page_37_Figure_1.jpeg)

#### **Result:**

influence on the correlated water motion of up to 20 Å, greater than the observed structural correlation length

This corresponds to 1000 H<sub>2</sub>O

S. Ebbinghaus, S. J. Kim, M. Heyden, X. Yu, U. Heugen, M. Gruebele, D. Leitner, and M. Havenith, PNAS 104, 20749 (2007)

### THz Spectroscopy of a prototype system: λ-repressor

![](_page_38_Picture_1.jpeg)

![](_page_38_Figure_2.jpeg)

structure (Protein + static hydration shell)

**Solvation dynamics** 

#### **Molecular modeling confirms:** Distance dependence of collective THz modes

![](_page_39_Figure_1.jpeg)

#### Dipole auto correlation function for different protein distances

![](_page_40_Figure_1.jpeg)

#### ∫dt exp(-i2Πt)<M(0)M(t)>

M is total dipole of system *n* is index of refraction THz dynamics of hydration water: Dependence on protein sequence and environment

![](_page_41_Picture_1.jpeg)

What about unfolding ? Does it influence the hydration dynamcis ?

![](_page_42_Figure_0.jpeg)

![](_page_42_Figure_1.jpeg)

Significant differences observed for the protein induced changes on the solvation water network dynamics when brought closer to unfolding: *S. Ebbinghaus, S. J. Kim, M. Heyden, X. Yu, U. Heugen, M. Gruebele, D. Leitner, and M. Havenith, JACS 101021 ja0746520 (2008)* 

# Comparison of the integrated THz absorbance (between 2.1 and 2.8 THz) of the pseudo-wild type lambda repressor with three mutants (Gln 22 →Tyr) of the protein

![](_page_43_Figure_1.jpeg)

Site directed mutagenesis:

replacement of polar glutamine sidechain by less polar sidechain when stabilizing secondary structures Gly4648Ala or destabilizing Ala3749Gly

# Long range water structuring if the proteins are closer to the unfolding transition

![](_page_44_Figure_1.jpeg)

Molecular dynamics simulation: retardation of water dynamics caused by exposure of hydrophobic residues to the water Is the long range water struturing important for the protein function ????

#### **Molecular dynamics simulation**

![](_page_45_Figure_1.jpeg)

Water molecules around the denatured state show retardation of dynamics caused by the exposure of hydrophobic residues of the denatured protein

#### X ray structure measures bound water THz spectroscopy probes changes in solvation dynamics

![](_page_46_Picture_1.jpeg)

Shown is tryptophan and bound water molecules (green dots)

Mapping of hydration dynamics Zhang; Wang; Kao; Qui; Yang; Okobiah; Zhong, *PNAS* **2007, 104, 18461 - 18466.** 

Distinguish: bound water, H-bond formation , 1. hydration shell Long range effect on water dynamics, extending several layers

#### Heterogeneous solvation dynamics: THz spectroscopy of ubiquitin B. Born, S.J. Kim, M. Gruebele, M. Havenith, 2008

![](_page_47_Figure_1.jpeg)

**Results:** 

Ubiquitin: dynamical hydration layer 10-15 Å

Introduction of tryptophan changes hydration dynamics !

 $\begin{array}{l} \text{mutation at surface:} \\ \alpha_{\text{solv}} \downarrow \text{(at 1.5 mM)} \\ \text{flexibility increases} \\ \alpha_{\text{solute}} \downarrow \text{(at 3.5 mM)} \end{array}$ 

Probing heterogeneous solvation dynamics:

**UB\*VA** and the **Ub\*AV** mutant show an increased flexibility due to the truncation of the sidechain at position 61; **Ile61Ala** replacement closer to the protein surface

#### THz spectroscopy of ubiquitin

- Wild type shows the most pronounced effect !
- Dynamical hydration layer exceeds to 10-17 Å Smaller than for λ-repressor
  - Replacement of phenylalanine by tryptophan
    influences hydration dynamics !

 Increased surface hydrophobicity of the mutants is a candidate for changes in hydration water yielding bulk water-like THz absorption

 largest deviation from Ub\* for the more disruptive core mutation Ala26Val, but a smaller effect for the less disruptive Ile61Ala mutation of a near surface

# Comparison of different experimental techniques to probe solvation dynamics:

![](_page_49_Figure_1.jpeg)

#### What about THz microscopy in a cell?

<u>Noninvasive</u> three-dimensional characterization of coupled network motions in complex heterogeneous systems in space <u>and</u> time (i.e. living cells and tissue)

What do we expect ?

Water in cells is not bulk like !

Water dynamics influence protein dynamics

And: Protein influences the water dynamics

**Biological relevance ?** 

![](_page_50_Picture_7.jpeg)

#### IR and THz spectroscopy: A tool to probe hydration dynamics

![](_page_51_Picture_1.jpeg)

![](_page_51_Picture_2.jpeg)

- Microsolvation can be probed by IR spectroscopy: (HCl)<sub>x</sub> (H<sub>2</sub>O)<sub>y</sub> in He nanodroplets, *J. Phys. Chem. A*, 111, 12192 (2007)
- THz spectroscopy probes the the dynamical size of the hydration layer (retardation of fast dynamics) for carbohydrates, *PNAS* 103, 12301 (2006)
- For proteins a non linear concentration dependence of the THz absorption indicates a long range influence on solvent dynamics of up to 20 Å; PNAS 104, 20749 (2007)
- Systematic study of denaturated proteins and site directed mutants of lamba repressor:

Probe of inhomogeous hydration dynamics, JACS 2008, in press

### Laser Spectroscopy & Biophotonics

#### (physicists, chemists, biochemists)

#### **Intermolecular Interaction**

![](_page_52_Picture_3.jpeg)

**Erik** Gerhard Bründermann Schwaab

![](_page_52_Picture_5.jpeg)

![](_page_52_Picture_6.jpeg)

Stephan Rudolph

Anna **Gutberlet** 

Anja Metzethin

Funds: THz **State: HBFG German science** foundation: **SP 1116 FOR 618** EU: **INTCHEM** International **HFSP Award 2004** 

![](_page_52_Picture_11.jpeg)

Konstanze **Matthias** Schröck Krüger

Simon

Udo **Ebbinghaus** Heugen

Markus

Ortlieb

**Benjamin** 

Born

Götz

Wollny

![](_page_52_Picture_16.jpeg)

**Matthias** Heyden

**Imaging: Scanning Nearfield** Infrared Microscopy

![](_page_52_Picture_20.jpeg)

Jean-Sebastian Samson

llona Kopf

![](_page_52_Picture_24.jpeg)

![](_page_53_Picture_0.jpeg)

### Classical Hole Motion in $\mathbf{E} \perp \mathbf{B}$

**Free hole motion under Lorentz-Force at low temperature (T = 4K)** 

$$\vec{\mathbf{F}} = \mathbf{m}^* \cdot \frac{\mathbf{d}\vec{\mathbf{v}}}{\mathbf{d}\mathbf{t}} = \mathbf{e} \cdot (\vec{\mathbf{E}} + \vec{\mathbf{v}} \times \vec{\mathbf{B}})$$

![](_page_54_Figure_3.jpeg)

#### The p-Ge laser: a tunable high power THz laser **Classical Hole Motion in E** $\perp$ **B** $\vec{\mathbf{F}} = \mathbf{m}^* \cdot \frac{\mathbf{d}\vec{\mathbf{v}}}{\mathbf{d}\mathbf{t}} = \mathbf{e} \cdot (\vec{\mathbf{E}} + \vec{\mathbf{v}} \times \vec{\mathbf{B}})$ **Optical** F **Phonon** Ehh E<sub>OP</sub> = 37 meV E<sub>lh</sub> m\*L m\*H $p_{L,H} = m_{L,H}^* E/B$ p<sub>L</sub> 2p<sub>L</sub> 2р<sub>н</sub> **p**<sub>H</sub> $\mathbf{p}_{L,H} = \mathbf{m}_{L,H}^* \mathbf{E}/\mathbf{B}$

#### THz difference absorption spectrometer for precise measurements of absorption coefficients

![](_page_56_Figure_1.jpeg)

![](_page_56_Picture_2.jpeg)

p-Ge laser: a unique THz source

![](_page_57_Figure_0.jpeg)

L. A. Reichertz, O. D. Dubon, G. Sirmain, E. Bründermann, W. L. Hansen, D. R. Chamberlin, A. M. Linhart, H.P. Röser, E. E. Haller, Phys. Rev. **B56**, 12069-12072 (1997)

E. Bründermann, H.P. Röser, A.V. Muravjov, S.G. Pavlov, V.N. Shastin, Infrared Phys. Technol. 1, 59-69 (1995)

2.5275

### THz spectroscopy as a tool to determine the size and the "nature" of the hydration shell in the sub-psec regime

Fit to a three component model:

 $\alpha_{total} =$ 

$$V_{sw}/V \alpha_{sw} + V_{solute}/V \alpha_{solute}$$
  
+ (1-V<sub>sw</sub>/V-V<sub>solute</sub>/V)  $\alpha_{bulk}$ 

 $V_{sw}/V$ : relative volume of the solvation water  $\alpha_{sw}$ : absorption coefficient of the solvation water  $V_{solute}/V$ : relative volume of the solute  $\alpha_{solute}$ : absorption coefficient of the solute

### Individual versus coherent motions

![](_page_59_Picture_1.jpeg)

Hydration water: The american way

#### THz spectroscopy of thin layers and films

- Measurement of small quantities of a probe
- Measurements on thin films, and liquid drops with high absorption coefficents
- Sensitive detection of thin layers required:
- Solution: attenuated reflection spectroscopy

![](_page_60_Figure_5.jpeg)

Exponentially decaying electric field Penetration depth: wavelength depend  $d=26 \ \mu m @ \lambda=100 \ \mu m$ 

### THz Cavity Enhanced Attenuated Total Reflection Spectroscopy

![](_page_61_Figure_1.jpeg)

### Increasing sensitivity:

Design of highly reflecting non absorbing mirrors in the THz R. Schiwon, G. Schwaab, E. Bründermann, M. Havenith App. Phys. L. 83(20), 4119, (2003)

- Wanted: no absorption, high bandwidth
- Gold: highly reflectant but non transparent
- Dielelectric mirrors for the THz region

Solution: Reflectivity <99% Alternating materials: spacing:  $\lambda/4$ distinct dielectric constant: silicon ( $\epsilon$ =11.7) (n=3.42) air ( $\epsilon$ =1.0) (n=1.0)

![](_page_62_Figure_6.jpeg)

![](_page_62_Figure_7.jpeg)

### Cavity Enhanced THz ATR Spectroscopy

![](_page_63_Figure_1.jpeg)

#### **Experimental results:**

R. Schiwon, G. Schwaab, E. Bründermann, M. Havenith Appl. Phys. Lett. 86, 201116 (2005)

- Lactalbumim hydrolysate: commercially available, degraded protein, found in many biological systems
- Probe dissolved and spread on surface
- Detection limit:

ATR: 23 nmol/mm<sup>2</sup>: 1500 nm (d<sub>penetration</sub>/17) corresponding to 1000 layers of free amino acids CEATR: 0.8 nmol/mm<sup>2</sup>: 50 nm (d<sub>penetration</sub>/520) corresponding to 33 layers of free amino acids yielding 15%±3% reduction

 New: THz absorption measurements of thin films and surfaces

#### Future potential for medical applications: Monitoring the influence of pharmaceuticals on a single cell level in real time

![](_page_65_Picture_1.jpeg)

**Chemical Science** 

### **Research highlights**

by the Royal Society, November 2004

Tracking drugs in single cells "A non-invasive method To quantify water in living cells is being pioneerd in Germany"

#### **Future improvements:**

Tomography improved lateral resolution