

Soft Matter and Functional Materials

User Facilities – Scientific Activities – Future Perspectives

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March 2011

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Research on Soft Matter and Functional Materials at the Helmholtz-Zentrum Berlin für Materialien und Energie

The Institute of Soft Matter and Functional Materials

Soft Matter Science is located at the interface between physics, chemistry, and biology, where novel and fascinating research areas are emerging and interdisciplinary approaches are required. Systems belonging to the field of soft matter range from biological macromolecules and their function in life science to industrial colloids that are produced in millions of tons. Investigations of the structure and dynamics of these systems provide a major challenge inasmuch as their typical sizes are located between the atomistic scale as e.g. in the case of proteins and the macroscopic scale going up to dimensions of a cell. The dynamic range is equally impressive since it goes from nanoseconds to hours. Scattering methods are uniquely suited to analyze soft matter systems in a comprehensive way. Thus, methods as e.g. small-angle neutron or x-ray scattering have become indispensable tools in the field of soft matter science.

The Institute of Soft Matter and Functional Materials (F-I2) at the Helmholtz-Zentrum Berlin für Materialien und Energie (HZB), founded in 2009, is devoted to the basic understanding and possible applications of colloidal and nanoscopic systems, of protein structure and function, and of complete cellular compartments. A main focus of our Institute is the combination of beamlines with dedicated user laboratories in which complementary methods such as light scattering and electron microscopy are provided for our users. Thus, difficult and sensitive samples can be prepared in our laboratories and directly analyzed at our beamlines. Serving a broad community of scientists from molecular biologists to industrial chemists we provide and develop

- top-class, dedicated beamlines
- complex experimental set-ups (neutrons, photons & more)
- dedicated laboratory infrastructure
- theoretical support (analytical modeling and computer simulations)

Based on our in-house research we allocate excellent beamlines for our users together with an advanced infra-structure. We provide all necessary complementary methods comprising both experimental facilities (user labs, advanced experimental set-up, etc) and offer additional support in theory (MD-simulations). An interdisciplinary team of scientists, PhD-students and technicians from physics, chemistry and biology is working on these topics and closely interacting with our users. In this way our institute provides an inspiring scientific environment for our 12 PhD students.

The development of new beamlines and beamline components is geared by in-house research that is devoted to the following topics:

- Structure and dynamics of proteins
- Analysis of soft- and bio-interfaces as well as thin films
- Synthesis and analysis of the structure and dynamics of colloidal suspensions
- Structural analysis of soft matter by a combination of small-angle scattering and microscopy including x-ray microscopy

Special emphasis is focused on a close collaboration with university groups, particularly from the Berlin area. This collaboration has led to the inauguration of the **Joint Berlin MX Laboratory** where research institutions from Berlin (HZB, Freie Universität Berlin (FU), Humboldt-Universität zu Berlin (HU), Max-Delbrück-Zentrum für Molekulare Medizin, Leibniz-Institut für Molekulare Pharmakologie) work together in the field of macromolecular crystallography. Three MX-beamlines located at the HZB are managed by this group. In many respects this joint lab has become the model for successful collaboration of a Helmholtz Zentrum with universities.

At present, the **Joint Laboratory for Structural Research** (JLSR) is set up in which the HZB, the HU, and the Institut für Kristallzüchtung closely collaborate in the field of structural analysis of soft and hard matter, in particular using microscopic methods. In this lab the HZB and the Institute of Physics of the HU run a high-resolution electron microscope and a cryogenic transmission electron microscope starting in April 2011.

Moreover, F-I2 has participated in the preparation of four applications of Berlin universities in the **excellence program** of the **Deutsche Forschungsgemeinschaft (DFG)**. We take part in the application of the proposal for the new Sonderforschungsbereich HIOS led by the Institute of Physics of the HU. For the time being, we are working in two **Schwerpunktprogramme** of the DFG (Intelligente Hydrogele, Kolloidverfahrenstechnik). Moreover, the group of Matthias Ballauff has transferred three research grants in the **DFG Normalverfahren** from the University of Bayreuth to the HZB. Together with international partners we acquired two research grants in the **FP7** of the EU, one on microfluidics and catalysis and another one on the development of the in situ diffraction system in the Macromolecular Crystallography Group. We are also partners in a grant for the development of nanofocusing optics for x-rays with the X-ray Microscopy Group. Furthermore, there are several cooperations with international laboratories and industrial partners.

An important point in the research strategy of F-I2 is the close collaboration with other Helmholtz partners and a close linkage to the Helmholtz Portfolio. Here we applied for a **Helmholtz Virtual Institute** devoted to **Multifunctional Materials for Biomedicine** (leading Helmholtz-center: Helmholtz-Zentrum Geesthacht Zentrum für Material- und Küstenforschung (HZG)). Moreover, we take part in the proposal for the **Helmholtz Portfolio** subject "**Technology and Medicine**".

This report gives an overview of the facilities provided with according statistical information and the inhouse research of the Institute F-I2 at the HZB. It covers the time span from July 2009 up to March 2011. The main lines of research at F-I2 will be presented together with recent scientific results. Moreover, the CVs of the leading scientists of the institute will be given.

Organization of the Institute

The Institute is subdivided into seven groups and one junior research group:

- Macromolecular Crystallography (Dr. Uwe Müller, Dr. Manfred S. Weiss)
- Biophysics (Dr. Thomas Hauß)
- Colloid Physics (Dr. Günter Goerigk, Dr. Daniel Clemens)
- Interfaces (Dr. Roland Steitz)
- Colloid Chemistry (Dr. Yan Lu)
- X-ray Microscopy (Priv. Doz. Dr. Gerd Schneider)
- Soft Matter Theory (Prof. Dr. Joachim Dzubiella)
- Junior Research Group Polymer Physics (Dr. Sebastian Seiffert)

The Institute is firmly rooted in the scientific environment of Berlin: Dr. J. Dzubiella is W2-S Professor at the HU, Dr. G. Schneider is Privatdozent at the same university (both Dept. Physics), Dr. S. Seiffert has the position of a junior researcher at the FU, Institute of Organic Chemistry, Dr. R. Steitz has a “Lehrauftrag” at the University of Technology of Berlin, Dr. M. Weiss and Dr. U. Mueller are both teaching with a “Lehrauftrag” at the FU, Institute of Chemistry and Biochemistry, and Dr. M. Ballauff is Professor (W3-S) at the Dept. Physics of the Humboldt Universität zu Berlin. Dr. G. Goerigk has a “Lehrauftrag” at the University of Paderborn.

At present (March 2011) the Institute has 49 coworkers that include
11 permanent scientists
15 postdocs
12 PhD-students
5 technicians

The output of the Institute is 52 publications in refereed journals in 2009 and 61 in 2010. This includes 18 papers in journals with an impact factor above 7, 64 in journals with impact between 3 and 7, and 31 in journal below 3.



Beamlines and Laboratories of the Institute

The institute runs **four neutron beamlines** on the Wannsee-site and **four at the synchrotron** in Adlershof. The following tables give a survey of the beamlines and the labs of F-I2 together with the group running the facility:

<i>Beamline</i>	<i>Instrument</i>	<i>Group</i>
V1	Membrane diffractometer	Biophysics
V6	Reflectometer	Interfaces
V16	SANS	Colloid Physics
V18	BioRef	Interfaces
BL 14.1	MX-beamline	Macromol. Crystallography
BL 14.2	MX-beamline	Macromol. Crystallography
BL 14.3	MX-beamline	Macromol. Crystallography
U41	X-ray microscope	X-ray microscopy
Electron beam writer	Electron beam writer	X-ray microscopy

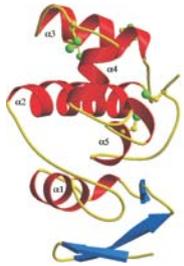
The Colloid Physics group collaborates with the Institute F-I1 in running the ASAXS-beamline at the synchrotron.

<i>Laboratories</i>	<i>Group</i>
BioLab	Macromol. Crystallography/Biophysics
Colloid Lab	Colloid Physics
Chemistry Lab	Colloid Chemistry
Laboratory for Microfluidics	Junior group Polymer Physics
Joint Berlin MX-Laboratory	Macromolecular Crystallography together with partners from the Berlin area
Joint Laboratory for Structural Research	Colloid Physics together with HU-Berlin

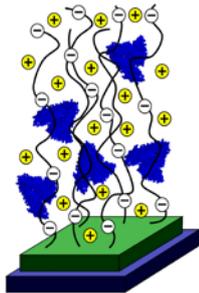
Soft Matter and Functional Materials: Research and Recent Highlights

The research of the institute is directed towards the analysis of soft matter throughout all the pertinent length scales as illustrated below. In particular, we study proteins and cellular components in general by starting on the atomistic level. Here a major effort of the Institute is devoted towards protein crystallography. The interaction of proteins with soft polymeric

Proteins



Bio-interfaces

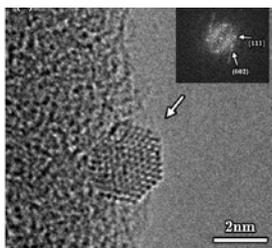


Cells

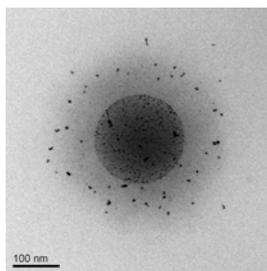


surfaces is a problem of high technical relevance. We investigate this interaction on planar surfaces as well as on nanoparticles by a wide range of methods as e.g. neutron reflection, SAXS, and calorimetry. Finally, entire cells can be studied by cryo-X-ray microscopy that allows us to visualize the different compartments of the cells without staining or microtome sectioning.

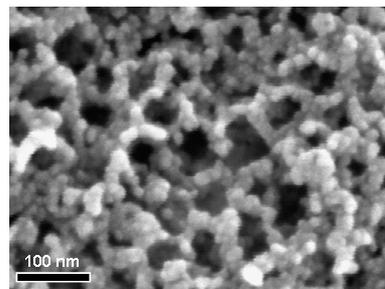
Nanoparticles



Composites



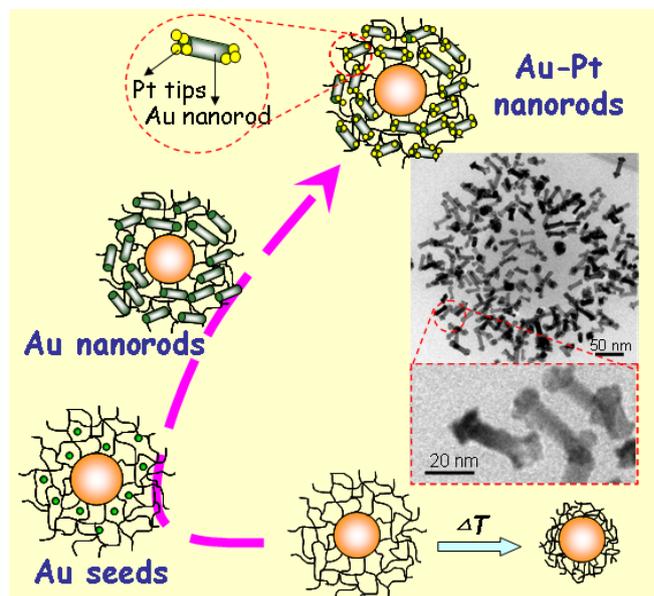
Mesoporous materials



Polymeric colloidal particles are studied in a similar way. Here we start from the synthesis of these systems including hybrids of inorganic nanoparticles with polymer colloids (see above). The various beamlines and labs provide an ideal place for analyzing these systems since all methods in the reciprocal and in direct space are available. The groups of the Institute work closely together in collaboration with many research institutions in the Berlin/Brandenburg area. In all cases we look into possible applications, again preferably together with strong partners. Therefore we also strengthen our collaborations with industrial partners in order to exploit possible applications as quickly as possible.

Polymers and nanocomposites

In-situ Growth of Catalytic Active Au-Pt Bimetallic Nanorods in Thermo-Responsive Core-Shell Microgels



We demonstrate that bimetallic Au-Pt nanorods (NRs) can be grown in situ into thermosensitive core-shell microgel particles by a novel two-step approach. In the first step, Au NRs with an average width of 6.6 ± 0.3 nm and length of 34.5 ± 5.2 nm (aspect ratio 5.2 ± 0.6) were homogeneously embedded into the shell of PNIPA networks. The volume transition of the microgel network leads to a strong red shift of the longitudinal plasmon band of the Au NRs. In the second step, platinum was preferentially deposited onto the tips of Au NRs to form dumbbell-shaped bimetallic

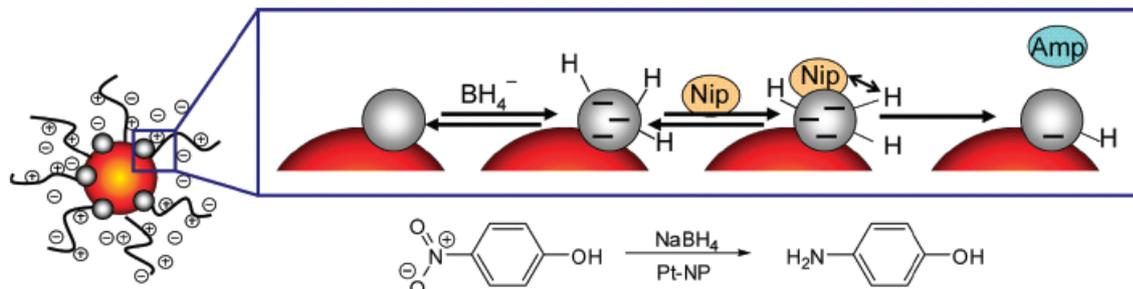
nanoparticles. The novel synthesis forms bimetallic Au-Pt NRs immobilized in microgels without impeding their colloidal stability. Quantitative analysis of the catalytic activity for the reduction of 4-nitrophenol indicates that bimetallic Au-Pt NRs show highly enhanced catalytic activity, which is due to the synergistic effect of bimetallic nanoparticles. The catalytic activity of immobilized Au-Pt NRs can be modulated by the volume transition of thermosensitive microgels. This demonstrates that core-shell microgels are capable of serving as “smart nanoreactors” for the catalytic active bimetallic nanoparticles with controlled morphology and high colloidal stability.¹

Kinetic Analysis of Catalytic Reduction of 4-Nitrophenol by Metallic Nanoparticles Immobilized in Spherical Polyelectrolyte Brushes

We present a study on the catalytic reduction of 4-nitrophenol by sodium borohydride in the presence of metal nanoparticles. The nanoparticles are embedded in spherical polyelectrolyte brushes, which consist of a polystyrene core onto which a dense layer of cationic polyelectrolyte brushes are grafted. The average size of the nanoparticles is approximately 2 nm. The kinetic data obtained by monitoring the reduction of 4-nitrophenol by UV/vis-spectroscopy could be explained in terms of the Langmuir-Hinshelwood model: The borohydride ions transfer a surface-hydrogen species in a reversible manner to the surface. Concomitantly, 4-nitrophenol is adsorbed and the rate-determining step consists of the reduction of nitrophenol by the surface-hydrogen species. The apparent reaction rate can therefore be related to the total surface S of the nanoparticles, to the kinetic constant k related to the rate-determining step, and to the adsorption constants $K(\text{Nip})$ and $K(\text{BH}_4)$ of nitrophenol and of borohydride, respectively. In all cases, an induction time $t(0)$ was observed of the order of minutes. The reciprocal induction time can be treated as a reaction rate that is directly related to the kinetics of the surface reaction because there is a linear

¹ Y. Lu, J. Yuan, F. Polzer, M. Drechsler, J. Preussner, *ACS Nano* **2010**, 4, 7078-7086.

relation between $1/(kt(0))$ and the concentration of nitrophenol in the solution. All data obtained for $t(0)$ so far and a comparison with data from literature indicate that the induction time is related to a slow surface reconstruction of the nanoparticles, the rate of which is directly related to the surface reaction.²



Mechanistic model (Langmuir-Hinshelwood mechanism) of the reduction of Nip by borohydride in the presence of metallic nanoparticles (gray spheres). The nanoparticles are bound to spherical polyelectrolyte brush (SPB) particles that consist of a polystyrene core and a shell of cationic polyelectrolyte chains. The catalytic reduction proceeds on the surface of the metal nanoparticles: The nanoparticles react with the borohydride ions to form the metal hydride. Concomitantly, nitrophenol adsorbs onto the metal surface. The adsorption/desorption of both reagents on the surface is fast and can be modeled in terms of a Langmuir isotherm. The rate-determining step is the reduction of the adsorbed Nip to Amp, which desorbs afterwards.²

Colloids and microemulsions

Adsorption of beta-Lactoglobulin on Spherical Polyelectrolyte Brushes: Direct Proof of Counterion Release by Isothermal Titration Calorimetry

The thermodynamics and the driving forces of the adsorption of beta-lactoglobulin on spherical polyelectrolyte brushes (SPB) are investigated by isothermal titration calorimetry (ITC). The SPB consist of a polystyrene core onto which long chains of poly(styrene sulfonate) are grafted. Adsorption isotherms are obtained from measurements by ITC. The analysis by ITC shows clearly that the adsorption process is solely driven by entropy while $\Delta H > 0$. This finding is in accordance with the proposed mechanism of counterion release: Patches of positive charges on the surface of the proteins become multivalent counterions of the polyelectrolyte chains, thereby releasing the counterions of the protein and the polyelectrolyte. A simple statistical-mechanical model fully corroborates the proposed mechanism. The present analysis shows clearly the fundamental importance of counterion release for protein adsorption on charged interfaces and charged polymeric layers.³

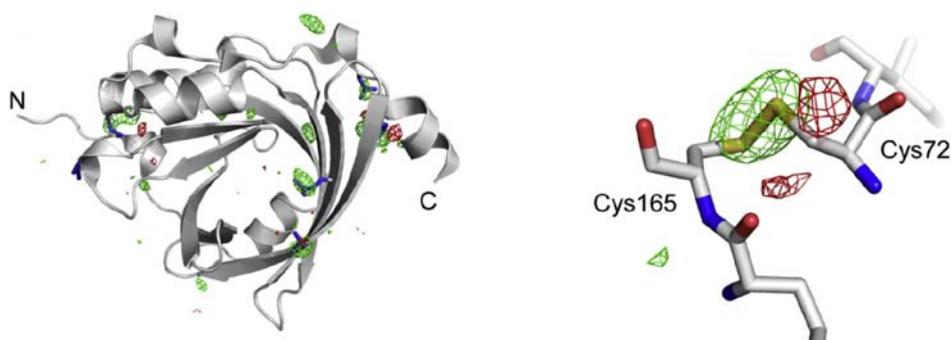
² S. Wunder, F. Polzer, Y. Lu, Y. Mei, M. Ballauff, *J. Phys. Chem. C* **2010**, *114*, 8814

³ K. Henzler, B. Haupt, K. Lauterbach, A. Wittemann, O. Borisov, Ballauff, M., *J. Am. Chem. Soc.* **2010**, *132*, 3159-3163

Cellular components

Phase determination Using the UV-Light Induced Radiation Damage

After the collection of an X-ray diffraction data-set from a macromolecule crystal the solution of the so-called “crystallographic phase problem” is the major task, which must be resolved. In order to achieve this, a growing number of methods exist, which we are aiming to extend with the further development of the UV-based radiation induced phasing (UVRIP) method. This experimental technique is focused on specific structural changes of cystine-containing protein crystals, which is due to the irradiation of the specimen with highly-intense UV-radiation. The structural changes can be used to work out a single isomorphous replacement (SIR)-like phasing scheme, which can lead to precise experimental phase information and thus to the access to the three dimensional structure (Figure). For this, a native data-set has to be collected before UV-exposure and compared with a second data set collected after the UV-irradiation of the same crystal. At the HZB-MX beamline BL14.1, we have installed all required instruments to carry out such experiments and are providing this to the user community. Within this research project, we aim to develop this method to reduce the existing requirements in terms of minimal data-set resolution and to investigate alternative specific damage sites within a macromolecular crystal.⁵



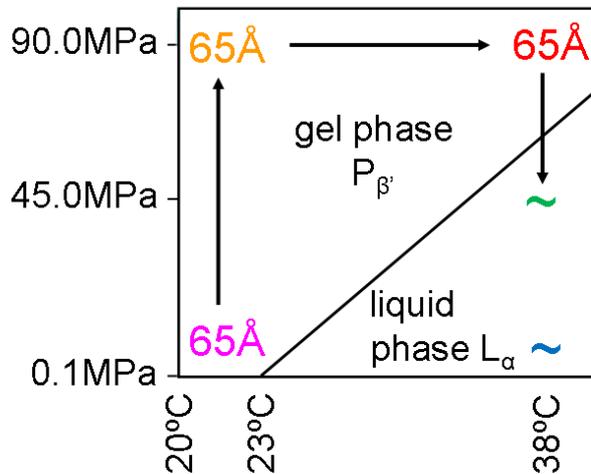
Specifically damaged disulfide-bridges of the protein α 1-acid glycoprotein

Pressure cell for Investigations of Solid-Liquid Interfaces by Neutron Reflectivity

We describe an apparatus for measuring scattering length density and structure of molecular layers at planar solid–liquid interfaces under high hydrostatic pressure conditions. The device is designed for in situ characterizations utilizing neutron reflectometry in the pressure range 0.1–100 MPa at temperatures between 5 and 60 °C. The pressure cell is constructed such that stratified molecular layers on crystalline substrates of silicon, quartz, or sapphire with a surface area of 28 cm² can be investigated against noncorrosive liquid phases. The large substrate surface area enables reflectivity to be measured down to 10⁻⁵ (without background correction) and thus facilitates determination of the scattering length density profile across

⁵ A. Faust, S. Puehringer, N. Darowski, S. Panjikar, K. Diederichs, U. Mueller and M. S. Weiss, *J. Appl. Cryst.* (2010). **43**, 1230-1237

the interface as a function of applied load. Our current interest is on the stability of oligolamellar lipid coatings on silicon surfaces against aqueous phases as a function of applied hydrostatic pressure and temperature but the device can also be employed to probe the structure of any other solid–liquid interface.⁶



Summary of experimental findings (d-spacings) for the oligolamellar DMPC bilayers film against excess water (D_2O), left. Note that the lipid film irreversibly detaches from support after crossing the phase boundary of the corresponding bulk system at 69 MPa and 38 °C. Photograph of the high pressure cell for neutron reflectometry (right).⁶

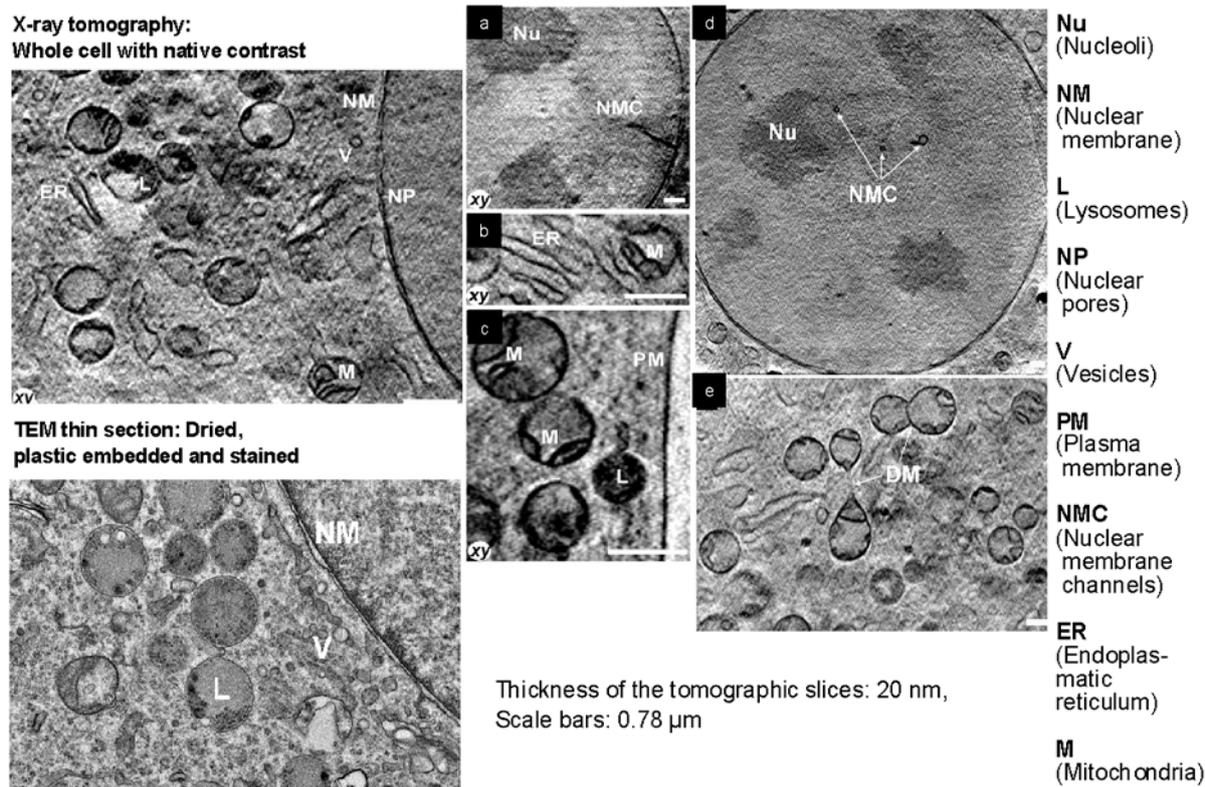
Biomaterials

Three-Dimensional Cellular Ultrastructure Resolved by X-ray Microscopy

We developed an X-ray microscope using partially coherent object illumination instead of previously used quasi-incoherent illumination. The design permitted the incorporation of a cryogenic tilt stage, enabling tomography of frozen-hydrated, intact adherent cells. We obtained three-dimensional reconstructions of mouse adenocarcinoma cells at 36 nm (Rayleigh) resolution, which allowed us to visualize the double nuclear membrane, nuclear pores, nuclear membrane channels, mitochondrial cristae and lysosomal inclusions.⁷

⁶ M. Kreuzer, T. Kaltfen, R. Steitz, B. H. Zehnder, R. Dahint, *Rev. Sci. Instr.* **2011**, *82*, 023902-7

⁷ D.L. Schönfeld, Ravelli R.B.G., Mueller U. and Skerra A., *J.Mol.Biol.* **2008** *384*, 393–405

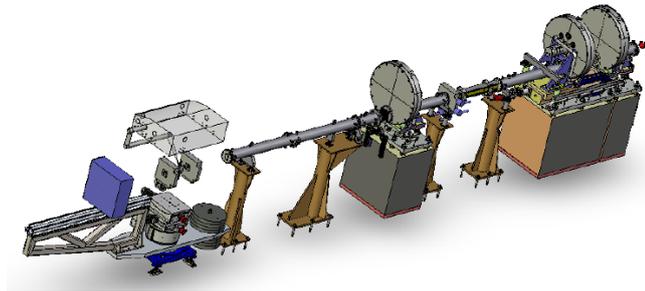
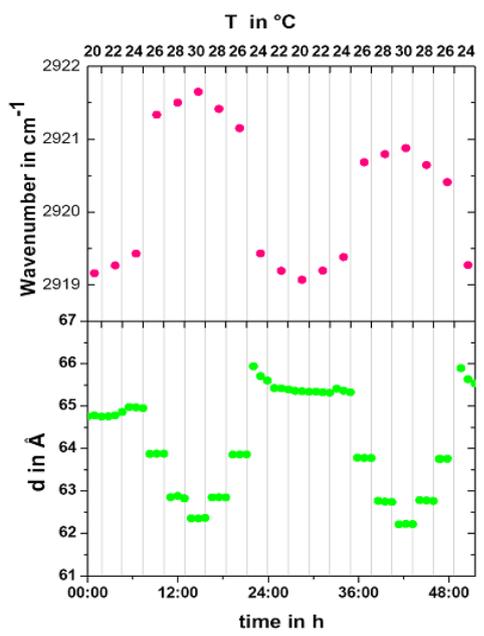


Comparison: Cryo X-ray tomography (upper left) and TEM thin section preparation (lower left). The slices of the X-ray tomograms (a-e) of frozen-hydrated mouse adenocarcinoma cells reveal numerous sub-cellular organelles including dividing mitochondria (DM), vesicles (V), the nuclear membrane (NM), nuclear pores (NP), nucleoli (Nu) and nuclear membrane channels (NMC).

BioRef – a Versatile Time-of-Flight Reflectometer for Soft Matter Applications at Helmholtz-Zentrum Berlin (in Cooperation with Ruprecht-Karls-Universität Heidelberg)

BioRef is a versatile novel time-of-flight (TOF) reflectometer featuring a sample environment for in-situ infrared spectroscopy at the reactor neutron source BER II of the Helmholtz Zentrum Berlin für Materialien und Energie (HZB). After two years of design and construction phase the instrument has recently undergone commissioning and is now available for specular and off-specular neutron reflectivity measurements. BioRef is especially dedicated to the investigation of soft matter systems and studies at the solid/liquid interface. Due to flexible resolution modes and variable addressable wavelength bands that allow for focusing onto a selected scattering vector range, BioRef enables a broad variety of surface and interface investigations and even kinetic studies with sub-second time resolution. The instrumental settings can be tailored to the specific requirements of a wide range of applications. The performance is demonstrated by several reference measurements, and the unique option of in-situ on-board infrared spectroscopy is illustrated by the example of a phase transition study in a lipid multilayer film.⁸

⁸ M. Strobl, R. Steitz, M. Kreuzer, M. Rose, H. Herrlich, F. Mezei, M. Grunze, R. Dahint, *Rev. Sci. Instrum.* **2011** in press



Analysis of lipid bilayers by the new BioRef beamline: d-spacing of an oligolamellar lipid bilayers coating on solid support against an excess water phase and corresponding ATR-FTIR signal of the asymmetric vibrational mode of CH₂ groups of the lipid chains as a function of sample temperature (left) as measured in-situ and combined at BioRef (right).⁸

Statistics User Service

List of Instruments run by the institute F-I2

Instruments in the BER II Cold Neutron Guide Halls

Instrument		Instrument scientist	Phone +49 30 8062-
V1	Membrane Diffractometer	Thomas Hauß	42071, 42202
V6	Reflectometer	Roland Steitz Ralf Köhler Robby Kischnik	42149, 42806 43077, 42806
V16	Very Small Angle Neutron Scattering (VSANS)	Daniel Clemens Karsten Vogtt Marcel Straschewski	42280, 43281 10-812 43022, 43281 42292
V18	Reflectometer for biological applications (Bio Ref)	Markus Strobl Martin Kreuzer Werner Graf	42490 43069 42829, 35835

Instruments run at BESSY II

Instrument		Instrument scientist	Phone +49 30 8062
BL 14.1	state-of-the-art MX beamline	Uwe Müller Karthik S. Paithankar	14974 15156
BL 14.2	beamline for de novo structure solution using anomalous phasing methods	Karthik S. Paithankar Sandra Pühringer	15156 15156
BL 14.3	fixed energy beamline	Manfred S. Weiss	13149
U41 X-ray Microscope	tomography of cryogenic samples	Peter Guttmann Stephan Werner Stefan Rehbein Gerd Schneider	14749 13181 13165 13131
Electron beam writer	VISTEC (type EBPG 5000+ ES)	Stefan Rehbein Stephan Werner Gerd Schneider	13165 13181 13131

Instrument Statistics 2009/I – 2011 / I
 (preliminary data, 2011, March)

V1: Membrane Diffractometer^{a)}

	2009/I	2009/II	2010/I	2010/II
number of accepted external proposals	5	6	4	4
number of allocated external n-days (short term)	47	52	52	39
cooperation (long term)	17	17	19	6
in-house EF proposals	2	2	3	1
in-house EF n-days	17	16	23	14
load factor external (<i>LFE</i>) ^{b)}	1.67	1.71	1.73	1.81

^{a)} Not in service for proposals due to neutron guide hall upgrade in 2011/I.

^{b)} *LFE* = number of requested days for experiments / remaining days for regular proposals

V6: Reflectometer

		Total number of proposals	Number of accepted proposals	Load factor external (<i>LFE</i>)
2009/I	Total	20	10	3.59
	short term extern	13	4	
	EF + LT	7	6	
	not scheduled		1	
	conducted		9	
2009/II	Total	20	11	2.46
	short term extern	13	6	
	EF + LT	7	5	
	not scheduled		0	
	conducted		11	
2010/I	Total	18	12	1.75
	short term extern	10	5	
	EF + LT ^{d)}	8	7	
	not scheduled		1	
	conducted		11	
2010/II	Total	19	7	5.05
	short term extern	14	3	
	EF + LT	5	4	
	not scheduled		1	
	conducted		6	
2011/I	Total	19	4	5.96
	short term extern	14	2	
	EF + LT	5	2	
	not scheduled		4	
	conducted		0 ^{c)}	

^{c)} Reactor shutdown 2010, Oct 2nd

^{d)} EF: in house research; LT: long term

V16: Very Small Angle Neutron Scattering (VSANS)

		Total number of proposals	Number of accepted proposals	Load factor external (LFE)
2010/II	Total	8	7	0.55
	short term extern	4	4	
	EF + LT	4	3	
	not scheduled		7	
	conducted		0	

Due to setup and commissioning, the only proposal session was for 2010/II.

V18: Reflectometer for biological applications (Bio Ref)

Until 2011/I, V18 was not available for public user service (proposals).

BL 14.1, BL 14.2, BL 14.3

	2009/I	2009/II	2010/I	2010/II	2011/I
Number of external Proposals	143	153	160	153	143
Number of inhouse proposals	11	15	22	25	25
Shifts requested	952	1101	887	1005	885
Shifts granted	630	719	578	612	700
Load factor	1.22	1.41	1.13	1.29	1.13

U41 X-ray microscope

		2009/I	2009/II	2010/I	2010/II	2011/I
Number of proposals applied for beamtime:	total	22	30	22	23	18
	there from inhouse proposals	1	4	2	3	1
	with approved beamtime	12	17	6	12	12
	there from inhouse proposals	1	4	1	2	1
Number of approved shifts for:	In-house users	24	28	22	42	28
	German users	36	27	22	22	38
	EU users	13	24	29	24	30
	Non-EU users	38	24	7	21	20

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Deputy Head

Dr. Nikoline Hansen
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Dr. Manfred Weiss

Dr. Nora Darowski

Dr. Karthik Paithankar

Michael Hellmig

Michael Steffien

Dr. Sandra Pühringer

Ronald Förster

Dr. Michael Krug

Dr. Martin Bommer

Dr. Monika Ühlein

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Dr. Karsten Voggt

Dr. Silvain Prévost

Christian Schneider

Miriam Siebenbürger

Christian Rabe

Dr. Beate Brüning

Dr. Ralf Stehle

Interfaces
Dr. Roland Steitz

Dr. Ralf Köhler

Dr. Markus Strobl

Martin Kreuzer

Holger Herrlich

Matthias Reinhardt

Colloidal Chemistry
Dr. Yan Lu

Julian Kaiser

Frank Polzer

Stefanie Wunder

Shuang Wu

Bin Dai

Dr. Martin Hoffmann

Theory
Prof. Dr. Joachim
Dzubiella

NN

NN

X-Ray Microscopy
PD Dr. Gerd Schneider

Dr. Peter Guttman

Dr. Stefan Rehbein

Katja Henzler

Dr. Stephan Werner

Basel Tarek

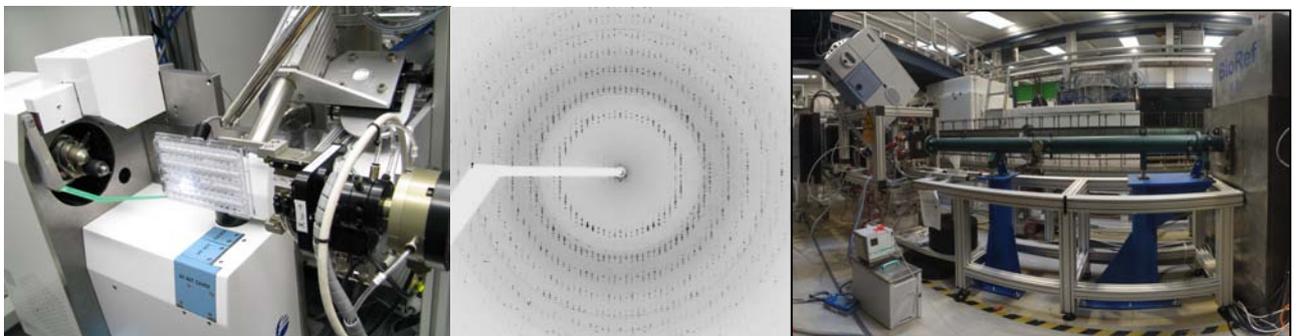
Polymer Physics
Dr. Sebastian
Seiffert

Torsten Rossow

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Soft Matter and Functional Materials

Beamlines



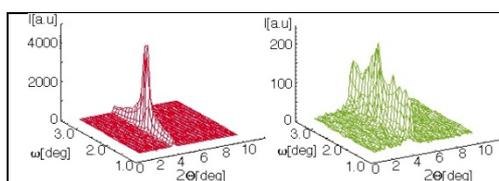
V1

Thomas Hauß

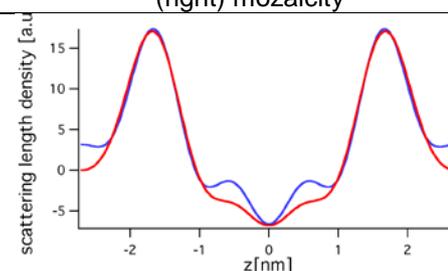


The diffractometer V1 with variable incident wavelength is installed at the curved neutron guide NL 1A. It is equipped with a high-resolution area detector. The design of the instrument is dedicated for experiments with biological membranes, polymers, micro-emulsions, micelles and other partly oriented systems. The high spatial resolution of the detector is appropriate for studying reflections from biological single crystals and magnetic satellite reflections.

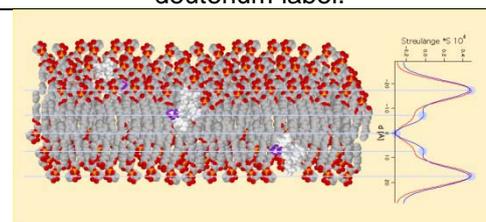
Type	Basic design: 2-axes diffractometer with cradle ($\pm 10^\circ$)
Monochromator	pyrolytic graphite (002), vertically focusing
Wavelength	selectable between 0.3-0.6nm (cold neutrons), corresponding to monochromator angles $2\Theta_M = 60^\circ$ - 120° (not alterable during experiment)
Angular Range	-10° to 120°
Collimation	$\gamma_0 = 1^\circ$ at 0.45 nm γ_1 : defined by two slit systems
Monochromator-Sample Distance	0.8m - 1.5m (extendable)
Sample-Detector Distance	0.8m - 2.0m
Detector	^3He , 19 x 19cm; pixel size 1.5 x 1.5mm ² ; height and inclination adjustable



Rocking curves around a Bragg peak of a membrane stack of low (left) and high (right) mozaicity



Neutron scattering length density profile of membranes with and without deuterium label.



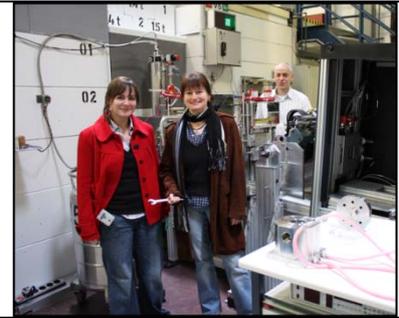
Localization of a specifically deuterated amyloid-b peptide in a lipid membrane with artistic model.

Selected publications

1. Schröter A, Kessner D, Kiselev MA, Hauß T, Dante S, Neubert RHH: *Basic nanostructure of stratum corneum lipid matrices based on ceramides [EOS] and [AP]: a neutron diffraction study*. Biophys J 2009, **97**(4):1104-1114.
2. Dante S, Hauß T, Brand A, Dencher NA: *Membrane fusogenic activity of the Alzheimer's peptide Aβ(1-42) demonstrated by small-angle neutron scattering*. Journal of Molecular Biology 2008, **376**(2):393-404.
3. Hauß T, Dante S, Haines TH, Dencher NA: *Localization of coenzyme Q₁₀ in the center of a deuterated lipid membrane by neutron diffraction*. Biochimica Et Biophysica Acta-Bioenergetics 2005, **1710**(1):57-62.
4. Dante S, Hauß T, Dencher NA: *β-amyloid 25 to 35 is intercalated in anionic and zwitterionic lipid membranes to different extents*. Biophysical Journal 2002, **83**(5):2610-2616.

V6

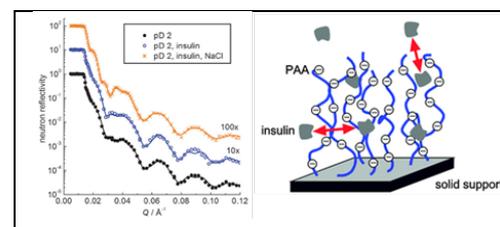
Roland Steitz
Ralf Köhler
Robby Kischnik



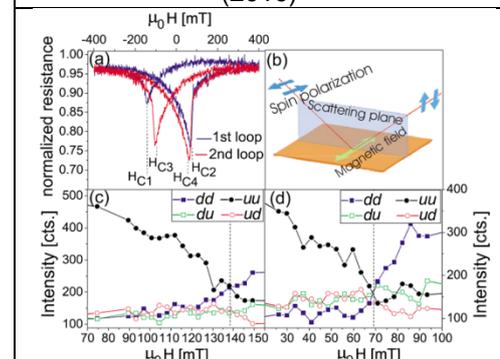
The reflectometer V6 is an angle dispersive fixed wavelength instrument dedicated to the investigation of thin films and surface structures at solid-air and solid-liquid interfaces as well as on free liquid surfaces.

The instrument is optionally equipped with polarization and polarization analysis for studies of magnetic thin films, also in external magnetic fields and at low sample temperature.

Monochromator	pyrolytic graphite (002) mosaicity: $\Delta\lambda/\lambda=2\%$
Wavelength	0.466 nm
Scattering plane	vertical
Polarization of neutron beam	98.5 %
Guide field	Permanent, horizontal
Detectors	^3He -detector tubes, position sensitive detector (180 x 180 mm, resolution 1.5 mm)
Q-range [$1/\text{\AA}$]	0 - 0.165 (0.127 free liquid surface)
Q-resolution [$1/\text{\AA}$]	0.001
Sample environment:	high pressure cell (1000 bar) for solid-liquid interfaces (RKU); heatable cells for liquids, solid-liquid and solid-gas interfaces; Langmuir Blodgett trough; horizontal magnetic field ≤ 1 T; sample rotation table (360°); closed cycle cryostat (4-300 K);



Probing adsorption and aggregation of insulin at a poly(acrylic acid) brush, Evers, F.; Reichhart, C.; Steitz, R.; Tolan, M.; Czeslik, C.; PCCP **12**, 4375 (2010)



Exchange bias by implantation of O ions into Co thin films, Demeter, J.; Meersschat, J.; Almeida, F.; Brems, S.; Van Haesendonck, C.; Teichert, A.; Steitz, R.; Temst, K.; Vantomme, A.; Appl. Phys. Lett. **96**, 132503 (2010)

Selected publications:

1. Interaction of IAPP and insulin with model interfaces studied using neutron reflectometry, Jeworrek, C.; Hollmann, O.; Steitz, R.; Winter, R.; Czeslik, C.; Biophysical Journal **96**, 1115 (2009)
2. Shear Induced Relaxation of Polymer Micelles at the Solid-Liquid Interface, Wolff, M.; Steitz, R.; Gutfreund, P.; Voss, N.; Gerth, S.; Walz, M.; Magerl, A.; Zabel, H.; Langmuir, **24**, 11331 (2008)
3. Binding of heavy and light water to polyelectrolyte multilayers, Ivanova, O.; Soltwedel, O.; Gopinadhan, M.; Koehler, R.; Steitz, R.; Helm, C. A.; Macromolecules, **41**, 7179 (2008)

V16

Daniel Clemens
Karsten Vogtt
Marcel Straschewski



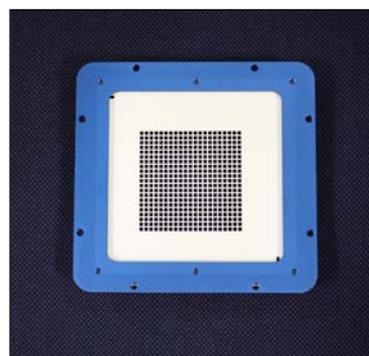
The very small-angle scattering instrument V16 (VSANS) serves for the analysis of mesoscopic structures, usually diluted in a buffer liquid. As the method is sensitive to variations in the scattering length density even pores can be investigated. Contrast variation techniques, namely deuteration of sections of the molecules to be investigated are crucial to obtain a maximum of information.

The sample table can be equipped with a thermalized 20 position sample changer, standard cryostats, furnaces or magnets.

Type	Basic design: TOF_SANS
Wavelength Band	0.2 - 1 nm (cold neutrons)
<i>Standard Mode:</i>	
Angular Range	0.2° to 30° (dependent on detector set-up)
Collimation	9 exchangeable guide optics: 1, 2, 4, 6, 8, 10 and 12m
Sample Detector Distance	1.7 m – 11.4 m
Q-range	$0.02 \text{ nm}^{-1} < Q < 16 \text{ nm}^{-1}$
Detector	$112 \text{ }^3\text{He-PSD}$ covering $100 \times 100 \text{ cm}^2$; pixel size $84 \times 84 \text{ mm}^2$
<i>Low-Q Mode:</i>	
Angular Range	0.05 to 0.75
Collimation	Multi-pinhole optics focusing on detector center
Sample-Detector Distance	11.4 m
Q-Range	$0.005 \text{ nm}^{-1} < Q < 0.4 \text{ nm}^{-1}$
Detector	^3He area covering $30 \times 30 \text{ cm}^2$; pixel size $2 \times 3 \text{ mm}^2$



20-Position sample changer, windows temperature controlled, removable



One of 24 Multi-pinhole diaphragms with 441 pinholes of $\sim 2 \times 2 \text{ mm}$

Selected publication:

1. Mezei, F., Clemens, D., Mokrani, L., *Neutronenoptisches Bauelement für die Neutronenkleinwinkelstreuungstechnik* - PCT/DE03/02869

V18

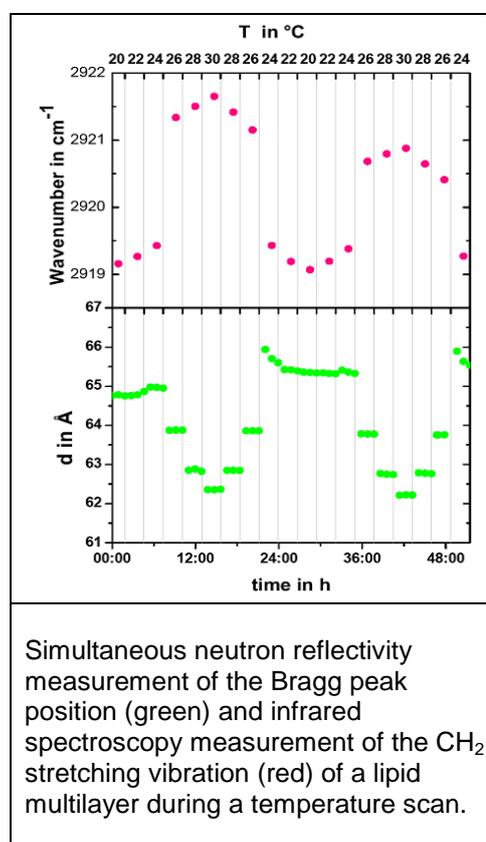
Markus Strobl
Martin Kreuzer
Werner Graf



The reflectometer V18 BioRef is a versatile time-of-flight instrument dedicated to the investigation of soft matter thin films and interfacial structures at solid-air and solid-liquid interfaces. The chopper system allows for tailoring the resolution of the instrument to the requirements of the specific measurement as well as for kinetic studies.

The instrument is optionally equipped with an infrared spectrometer for simultaneous in-situ measurements in ATR-FTIR geometry.

3-chopper system	$\Delta\lambda/\lambda=1\%$ to 12%
Wavelength band	0.25 nm to 0.7 - 1.8 nm
Scattering plane	horizontal
Polarization of neutron beam	Not yet
Guide field	Not yet
Detectors	position sensitive detector (300 x 300 mm, resolution 2 x 3 mm ²)
Q-range [1/Å]	0 - 0.4
Q-resolution [dQ/Q]	1.4 – 20%
<i>Sample environment:</i>	
Heatable cells for solid-liquid and solid-gas interfaces; In-situ ATR-FTIR spectroscopy Pressure and shear environment under development	



Selected publications:

1. M. Strobl, R. Steitz, M. Kreuzer, A. Nawara, F. Mezei, M. Rose, M. Grunze and R. Dahint, *BioRef – a time-of-flight neutron reflectometer combined with an in-situ infrared spectrometer at the Helmholtz Centre Berlin*, J. of Phys. (conference series) **251** (2010) 012059
2. M. Strobl, R. Steitz, M. Kreuzer, M. Rose, H. Herrlich, F. Mezei, M. Grunze, R. Dahint, *BioRef – a versatile time-of-flight reflectometer for soft matter applications at Helmholtz-Zentrum Berlin für Materialien und Energie*, Berlin, Rev. Phys. Instrum. (2011)

BL14.1

Uwe Müller



BL14.1 is a state-of-the-art MX beamline and currently the most modern and efficient MX beamline in Germany. It is energy-tunable within the range from 5 keV (2.5 Å) to 16.5 keV (0.75 Å). BL14.1 is equipped with an automatic sample handling robot (CATS). The MD2 microdiffractometer, which is equipped with a mini-kappa goniometer and on-axis sample zoom-microscope, enables the visualization and 3D-centring of crystals at a micrometer scale in the X-ray beam.

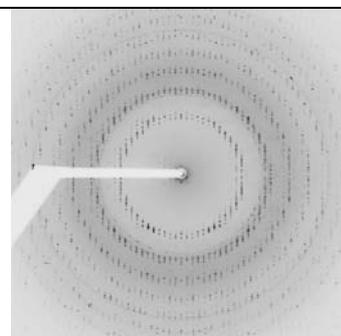
Energy range [keV]	5.5 -15.5
Wavelength range[Å]	0.80-2.25 (max. intensity at 0.92)
Max. photon flux at sample [Phot/s/0.1A/0.05% BW]	1.3×10^{11} (13 keV)
Energy resolution [eV]	< 2 (9 keV)
Goniometry	Microdiffractometer with Mini-kappa
Sample automation	CATS sample mounting robot (handling of up to 90 SPINE compatible samples)
X-ray detector	Rayonics MX-225
Beam size [μm]	30-100 diameter
Achievable resolution [Å]	0.9
Maximum cell parameter [Å, at 2.0 Å resolution]	400
Exposure time range [sec]	1 - 20

Experimental possibilities:

- High performance *de novo* structure determination by MAD, SAD, SIRAS, MIRAS
- Efficient Screening
- Handling of very small crystals
- RIP, UVRIP
- Element identification using X-ray fluorescence



11 μm crystal within a 50 μm X-ray beam



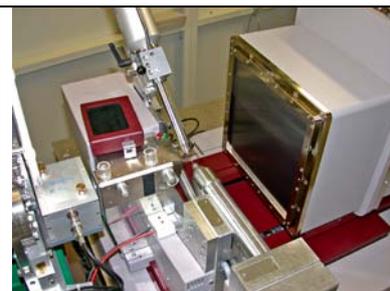
Diffraction image of an aligned protein crystal with reciprocal lattice vector along the X-ray beam

Selected publications:

1. Gao S; Malsburg A; Paeschke S; Behlke J; Haller O; Kochs G; Daumke O, *Structural basis of oligomerization in the stalk region of dynamin-like MxA* 2010, Nature 465, 7297
2. Luckner, S.R., Machutta, C.A., Tonge, P.J., Kisker, C., *Crystal Structures of Mycobacterium Tuberculosis Kasa Show Mode of Action within Cell Wall Biosynthesis and its Inhibition by Thiolactomycin* 2009, Structure 17, pp. 1004
3. Monecke, T., Guttler, T., Neumann, P., Dickmanns, A., Gorlich, D., Ficner, R., *Crystal Structure of the Nuclear Export Receptor CRM1 in Complex with Snurportin1 and RanGTP* 2009, Science 1087

BL14.2

Karthik S.
Paithankar
Sandra Pühringer

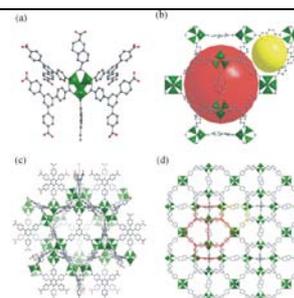


BL14.2 is the workhorse beamline for *de novo* structure solution using anomalous phasing methods such as MAD. The beamline is energy-tunable within the range from 5 keV (2.5 Å) to 16.5 keV (0.75 Å). BL14.2 is equipped with a mardtb goniometer, which makes it possible to achieve very short detector-to-sample distances of down to 45 mm.

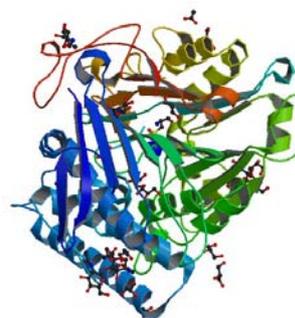
Energy range [keV]	5.5 -15.5
Wavelength range [Å]	0.80-2.25 (max. intensity at 0.92)
Max. photon flux at sample [Phot/s/0.1A/0.05% BW]	1.9×10^{11} (13 keV)
Energy resolution [eV]	<2 (9 keV)
Goniometry	MARdtb
Sample automation	No
X-ray detector	Rayonics MX-225
Beam size [µm]	100 x 150

Experimental possibilities:

- High performance *de novo* structure determination by MAD, SAD, SIRAS, MIRAS
- Long wavelength phasing like S-SAD
- Small molecule crystallography application
- Ultra-high resolution data collection
- Element identification using X-ray fluorescence



Crystal structure of mesoporous metal-organic framework (Klein et al. 2009)



Crystal structure of a 66.3 kDa protein solved by S-SAD (Lakomek et al. 2009)

Selected publications:

1. I Grueninger, D., Treiber, N., Ziegler, M.O.P., Koetter, J.W.A., Schulze, M.-S., Schulz, G.E., *Designed Protein-Protein Association* 2008, Science 319, 206
2. Klein, N., Senkovska I., Gedrich U, Stoeck U, Henschel A, Mueller U and Stefan Kaskel, *Eine mesoporöse Metall-organische Gerüstverbindung* 2009, Angew. Chem. Int. Ed. Vol. 48, 9954-9957
3. Lakomek, K., Dickmanns, A., Mueller, U., Kollmann, K., Deuschl, F., Berndt, A., Lubke, T., Ficner, R., *De novo sulfur SAD phasing of the lysosomal 66.3 kDa protein from mouse* 2009, Acta Crystallogr., Sect.D 65, 220

BL14.3 Manfred S. Weiss



BL14.3 is a fixed energy beamline which is operated at an energy of 13.8 keV (0.89 Å). It can be used for *de novo* structure solution using anomalous phasing methods such as SAD utilizing heavy atoms like Pt, Hg, Au, Se and others. BL14.3 offers a unique experimental set-up to improve the diffraction properties of protein crystals. It is equipped with a HC1c dehydration setup, which can be used to improve the diffraction quality of crystals by controlled dehydration.

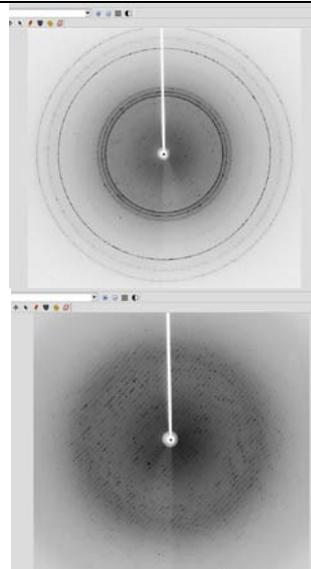
Energy [keV]	13.8
Wavelengths [Å]	0.89
Max. photon flux at sample [Phot/s/0.1A/0.05% BW]	4×10^{10} (13.8 keV)
Energy resolution [eV]	< 5
Goniometry	MARdtb
Sample automation	n/a
X-ray detector	Rayonics SX-165
Beam size [µm]	100 x 200
Achievable resolution [Å]	0.9
Maximum cell parameter [Å, at 2.0 Å resolution]	250
Average exposure time [sec]	3 - 30

Experimental possibilities:

- High performance *de novo* structure determination by SAD, SIRAS, MIRAS
- Crystal annealing with a remotely controlled cryo-shutter
- Controlled dehydration of protein crystals
- High resolution data collection



BL14.3 annealing device in operation



Macromolecular crystal diffraction image before and after annealing

Selected publication:

Kuettner, E.B., Kettner, K., Keim, A., Svergun, D.I., Volke, D., Singer, D., Hoffmann, R., Muller, E.C., Otto, A., Kriegel, T.M., Straeter, N., Crystal Structure of Hexokinase KIHxk1 of *Kluyveromyces lactis*: A MOLECULAR BASIS FOR UNDERSTANDING THE CONTROL OF YEAST HEXOKINASE FUNCTIONS VIA COVALENT MODIFICATION AND OLIGOMERIZATION (2010) *J.Biol.Chem.* **285**, pp. 41019-41033

U41 X-ray Microscope

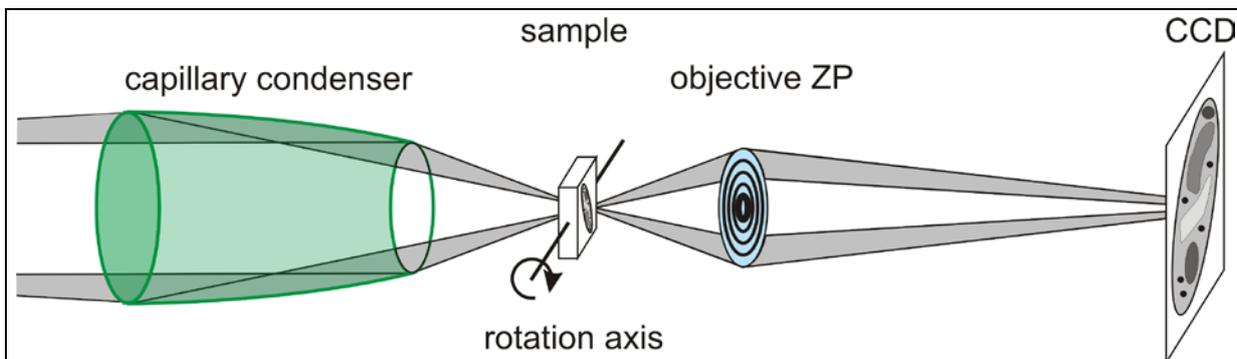
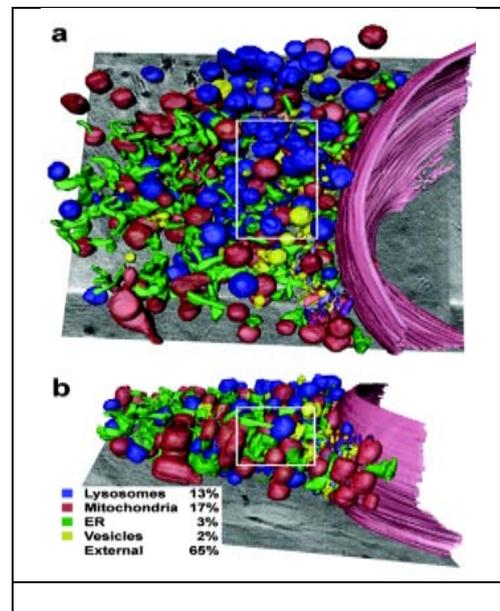
Peter Guttman
Stephan Werner
Stefan Rehbein
Gerd Schneider



The X-ray microscopy group at the Helmholtz-Zentrum Berlin specializes in the development and application of advanced X-ray microscopy, X-ray tomography and X-ray optics for the 10-nm scale characterization of the nanostructure, chemical nature, and composition of materials with high energy resolution. This world-class facility is comprised of a unique combination of state-of-the-art X-ray microscopy instruments, in-house X-ray diffractive optical development and staff members with expertise in microscopy, physics, biophysics and chemistry to address national needs and technical challenges that impact materials, energy and life sciences.

The full-field cryo transmission X-ray microscope provides unique capabilities for high resolution X-ray imaging. It permits tomography of cryogenic samples on flat sample holders as well as spectromicroscopy studies with high energy resolution $\Delta E/E=10^{-4}$ at nanoscale lateral resolution.

Type	Cryo full-field transmission X-ray microscope
Photon energy range	0.25 – 1.5 keV
Energy resolution	10^{-4}
Sample temperature	- 170° C - room temperature
Sample tilt	$\pm 80^\circ$
X-ray source	U41 undulator
3D spatial resolution	25 nm
2D spatial resolution	10 nm
Monochromator	SGM



Selected publications:

1. S. Rehbein, S. Heim, P. Guttman, S. Werner, G. Schneider, *Ultra-high-resolution soft-x-ray microscopy with zone plates in high orders of diffraction*, Phys. Rev. Lett. **103**, (2009) 110801
2. G. Schneider, P. Guttman, S. Heim, S. Rehbein, F. Mueller, K. Nagashima, J.B. Heymann, W.G. Müller, J.G. McNally, *Three-dimensional cellular ultrastructure resolved by X-ray microscopy* Nature Methods **7** (2010), 985-987
3. S. Heim, P. Guttman, S. Rehbein, S. Werner, G. Schneider, *Energy-tunable full-field x-ray microscopy: Cryo-tomography and nano-spectroscopy with the new BESSY TXM*, Journal of Physics: Conference Series **186** (2009) 012041

Electron Beam Writer

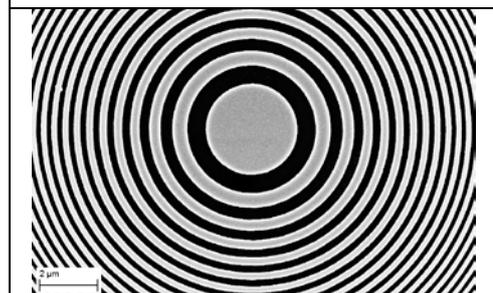
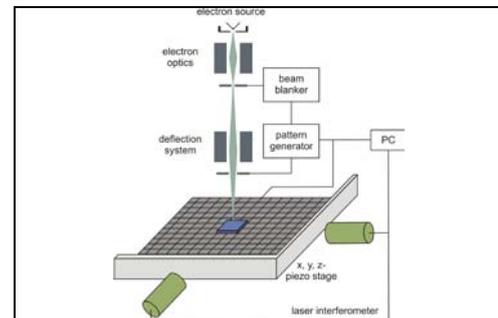
Stefan Rehbein
Stephan Werner
Gerd Schneider



The X-ray microscopy group in the institute Soft Matter and Functional Materials operates a state-of-the-art electron beam writer from VISTEC (type EBP 5000+ ES). With its small electron beam size in the range of few nanometers and the high electron energy of 100 keV, lithography with nanometer precision is possible. Arbitrary pattern can be computer generated, converted into machine readable format and finally exposed. The writing field size without moving the wafer stage is 256 μm at 100 keV electron energy. Larger areas can be exposed by stitching the writing fields using the laser interferometer controlled wafer stage. Under these conditions maximum areas of 4 inch can be exposed, depending on the writing time which increases with smaller step sizes. Another advantage of our e-beam tool is the high overlay precision in the 10 nm range which permits to stack processed layers. The application fields for the new e-beam system are diffractive X-ray optical elements, for example high-resolution Fresnel zone plates for X-ray microscopy or advanced monochromator gratings.

Specifications of the VISTEC EBP 5000+ ES

Electron energy	50, 100 keV
Writing field size	256 μm (100 keV)
Beam current	100 pA – 100 nA
Pattern generator frequency	25 MHz
DAC	16 Bit (main-field), 14 Bit (sub-field)
Spot size	2.2 nm (100 keV)
On-axis resolution in resist	8 nm
Laser stage resolution	0.6 nm
Wafer size	up to 4 inch



Selected publications:

1. S. Rehbein, S. Heim, P. Guttman, S. Werner, G. Schneider, *Ultra-high-resolution soft-x-ray microscopy with zone plates in high orders of diffraction*, Phys. Rev. Lett. **103**, (2009) 110801
2. S. Werner, S. Rehbein, P. Guttman, S. Heim, G. Schneider, *Towards high diffraction efficiency zone plates for X-ray microscopy* Microelectron. Eng. **87** (2010), 1557-1560
3. S. Rehbein, S. Werner, P. Guttman, S. Heim and G. Schneider, *Soft X-Ray Microscopy at HZB: Zone Plate Development and Imaging using the 3rd Order of Diffraction*, 10th international conference on x-ray microscopy, AIP Conference Proceedings (2010)

Soft Matter and Functional Materials

Laboratories



The BioLab: Thomas Hauß, Manfred Weiss

The BioLab is an essential service unit for the HZB user platform and provides on-site sample preparation and characterisation in parallel and complementary to neutron and X-ray scattering experiments. The Wannsee branch of the BioLab is run by the Biophysics group and it is routinely used by neutron users and cooperation partners with the scientific background soft matter and biology, a community of 1/3rd of all neutron users. The BioLab offers biophysical, biochemical, and cell laboratories with a broad range of laboratory-based equipment. Specialised sample environments for neutron scattering experiments, especially complementary to neutron diffraction and reflectometry, SANS, SAXS, INS and QENS are developed. The expertise ranges from membrane biophysics and structural biology over protein dynamics to structure and function of interfaces. Significant achievements for the user support are sophisticated preparation methods for model membranes, a reliable and robust protocol for the preparation of “free floating bilayers” for neutron reflectivity, and a novel real-time (laser-neutron) pump-probe experiment to study modulation in protein dynamics by neutron scattering methods. Future developments will strengthen experiments under most physiological condition, establish in-situ sample characterisation like in-situ ATR-FTIR spectroscopy, and offer new characterisation methods like dynamic light-scattering and fluorescence microscopy.

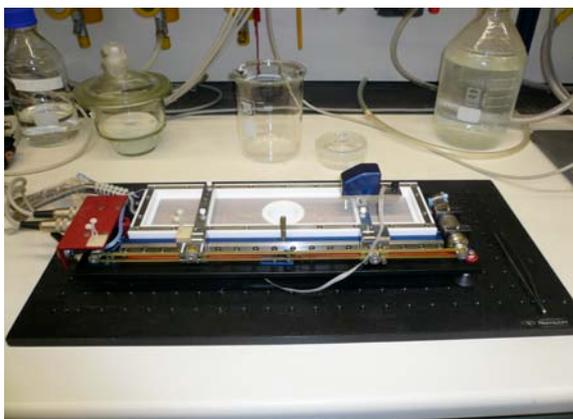
The Adlershof branch of the BioLab is run by the MX-group and is currently being converted into a protein production facility for Structural Biology experiments. Once completed, it will be possible to perform all steps from cloning, heterologous bacterial expression, protein purification and characterisation as well as crystallization of proteins for X-ray diffraction experiments. The BioLab will support the research activities of the HZB MX-group but it will also remain available to other groups as well as external users.



Left: Fluorescence microscope with spot illumination



Right: Cell Laboratory



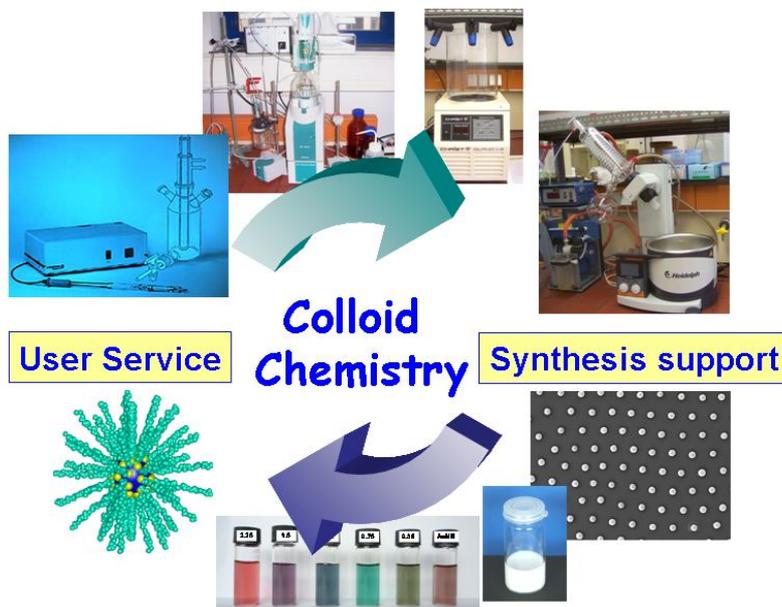
Langmuir trough



Laser laboratory

The Chemistry Lab: Yan Lu

The chemistry lab has all the facilities for the synthesis and the characterization of colloidal suspensions, micelles, and supra-molecular structures generated in aqueous suspension. The main task of chemistry lab is to give support for the users during their beam time in HZB. This includes not only supporting the user with full-equipped chemistry lab for sample preparation, but also giving them professional guide for synthesis of different colloidal particles (from organic to inorganic particles).



All together we run two synthesis labs (L130, L131), one sample purification lab (LS123), one sample preparation lab outside Neutron Halle (V108) and one user lab in Neutron Halle (UYH0343), which can be accessed by our users. All the synthesis labs and sample preparation labs are equipped with fume hoods. Supplements, such as Millipore water, magnetic stirrer, thermostat, vortex mixer, balance and research pipettes, etc. are available in all labs. Storing samples either in refrigerator/freezer ($\sim 10^{\circ}\text{C}$) or ultra-low temperature freezer ($\sim -20^{\circ}\text{C}$) is possible during beamtime.

Giving technique supports for synthesis is another core-task for the chemistry lab. Concerning the requests of the users, polymerization (such as emulsion polymerization, dispersion polymerization and Atom Transfer Radical Polymerization (ATRP)) can be conducted in the colloid chemistry lab. In addition, two UV-reactors are available in the synthesis lab that photo polymerization/reaction can be also carried out as requested. We supply the service of basic characterization of the colloidal samples as well (such as pH/conductivity measurement, auto-titration, zeta-potential etc.). Special sample treatment with high vacuum oven or freeze-drying is possible. In addition, this lab provides support by doing electron microscopic studies on colloids.

The combination of user service and professional technical support based on colloid chemistry provides an important research concept in the Institute of Soft Matter and Functional Materials.



The Colloid Lab: Characterization of particles with Light and Rheology:

Daniel Clemens



Dynamic light scattering instrument, manufactured by ALV

The colloid lab is run by the colloid physics group within the Institute for Soft Matter and Functional Materials. It encompasses laboratory methods that complement the work with the instruments on HZB's large scale facilities, which comprise thermally controlled static and dynamic light scattering (SLS/DLS). Moreover, the DLS instrument allows us to do depolarized dynamic light scattering as

well that gives highly valuable information about anisometric objects. Both DLS-instruments are situated in

our large class IIIb laser laboratory (LS117) together with a commercial Malvern Zeta-Sizer. Additionally we run three rheometers in a dedicated lab (LS217), one of which is foreseen to be adapted to the small-angle neutron scattering instrument V16. The group research on mesoscopic materials as colloidal suspensions, micelles and supramolecular structures is in this way fully cross-linked with the development of instruments as well as the accessibility to and service of supporting laboratory equipment.

Our equipment is available for guest groups, especially for our users that have been granted beam time on the neutron or synchrotron instruments, to allow for a complete and proper characterization of their samples. We plan to extend these facilities to establish more specialized sample environment as e.g. a rheo-SANS device for the VSANS-beamline. With these facilities we are near to a complete set of tools for the characterization of colloidal suspensions, micelles, and supramolecular structures generated in aqueous suspension.



Malvern Zeta-Sizer



Modular Anton Paar Physica MCR 301 rheometer for diverse geometries

Joint Laboratory for Structural Research

Soft matter systems consist of structures and structural units with sizes that span from the atomistic range to micrometers. Analysis of such systems hence requires a wide range of methods that have access to this range. Moreover, probes used for the structural analysis should be sensitive to the details of the system under consideration. The newly founded Joint Laboratory for Structural Research meets these demands and offers a wide range of methods summarized in the following table:

Partners	Groups	Methods
HU	Professorship for electron microscopy and structural research (W3)	HR-TEM of hybrid systems
	Prof. J.P. Rabe, PD S. Kirstein, Institut für Physik	AFM
	Prof. S. Kowarik, Institut für Physik	X-ray scattering and reflectivity
HZB	PD Dr. G. Schneider, Dr. K. Henzler, Soft Matter and Functional Materials	X-ray microscopy, E-beam lithography
	Prof. M. Ballauff, Dr. Y. Lu, Soft Matter and Functional Materials	Cryo-TEM
	Dr. G. Goerigk, Soft Matter and Functional Materials	SAXS, SANS
TU	Prof. Regine von Klitzing	Cryo-TEM

The new **cryo-TEM** of the JLSR will be set up in April 2011 and serve the community of soft matter scientists on the Adlerhof Campus and the users of the HZB. Its special features are:



- Variable acceleration voltage up to 200kV
- LaB6 cathode with cool beam illumination
- Minimum dose system for sensitive samples
- Alpha selector for illumination conditions
- Dedicated vacuum system for cryoTEM
- TVIPS 4x4k fast scan CCD camera
- Single tilt cryo holder (high tilt from 0° - 80°)

The Joint Berlin MX Laboratory: Uwe Müller, Manfred S. Weiss

Over the past years, Berlin has become the German center of X-ray crystallography based structural biology, with currently 12 independently operating macromolecular crystallography groups.

Within this group of researchers, the HZB-MX beamlines are a natural condensation point for most of the experimental work carried out within the 12 groups. Even more, the proximity of the MX-beamlines to the laboratories of the groups may even be the major reason for the success of the groups.

Consequently, in 2008 a more formal platform for collaboration has been established by the following institutes:

- Helmholtz Zentrum Berlin für Materialien und Energie
- Freie Universität Berlin
- Humboldt Universität zu Berlin
- Max-Delbrück-Zentrum für Molekulare Medizin in Berlin-Buch
- Leibniz-Institut für Molekulare Pharmakologie

These five institutes have founded the “Joint Berlin MX-Laboratory” as a new model for a multi-institutional research collaboration in Berlin.

Within the next four years the participating will work on a number of already initiated research projects, which were motivated by the collaboration partners and thus directly link the distinct research fields with each other. If possible the projects will be connected with each other, using the new possibilities available within this joint activity.

In addition to the research, the partners share the responsibility for the operation and future development of BL14.2-3 as well. First fund raising results could be achieved already through the submission of three joint research proposals, which did succeed in one BMBF-funded project, for the duration of three years. Within this project, the Freie Universität Berlin, HZB and the Russian Academy of science will work closely together within the field of structural biology of ribonucleoprotein complexes.



Figure 1: Inauguration of the Joint Berlin MX-Laboratory



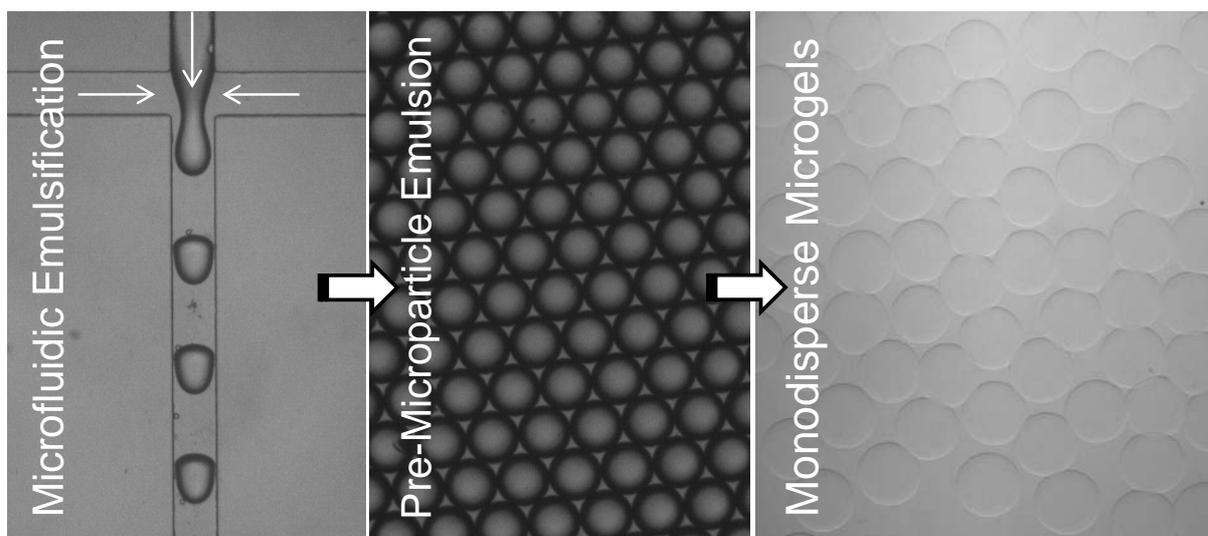
Figure 2: Meeting of 73 Berlin structural biologists at BESSY II during the 1. Joint MX-day in August 2010

Laboratory for Microfluidics: Sebastian Seiffert

Microfluidic devices are networks of micron scale channels integrated to perform functions. These functions can be divided into two broad classes. In one class, microfluidic devices perform chemical and biological assays by introducing cells, beads, and other reagents into the device and then merge, mix, and split them. In the second class, microfluidic devices fabricate fluid droplets of precisely controlled geometry, which then serve to synthesize microparticles. For this purpose, solutions or melts of monomers or crosslinkable polymers are introduced into the device, along with an immiscible carrier phase; the devices disperse these solutions into equally sized micro-droplets, which can then be solidified by polymerization, crosslinking, or crystallization, as illustrated in the figure.

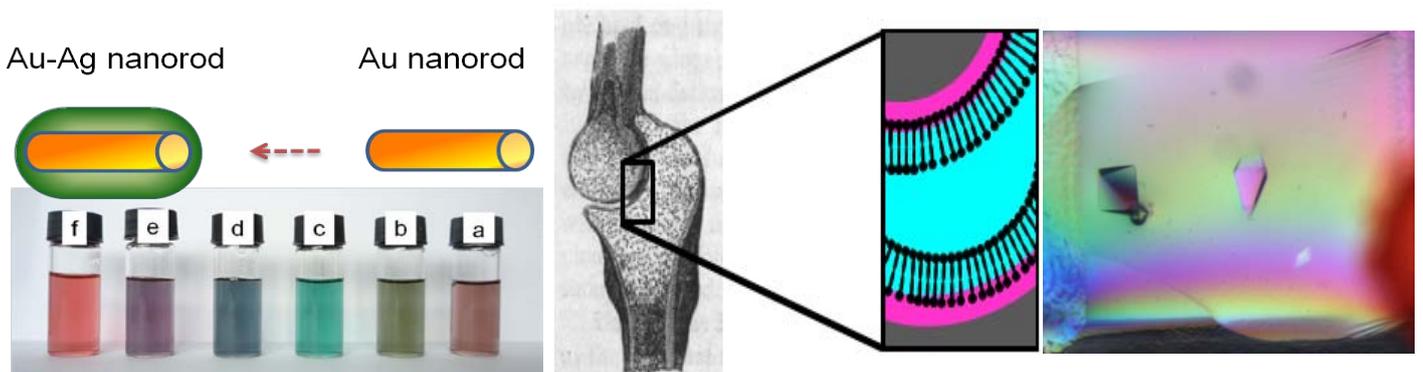
The principle of the microfluidic drop formation can be explained using a water faucet as an example: if a faucet operates a low flow rate, water drips out one drop at a time. The drop size is determined by a balance between the surface forces of the hanging drop and its weight, and therefore depends on the surface tension of the fluid and the size of the faucet. Since both these parameters are constant, all drops exhibit a narrow size distribution. The same principle is employed in microfluidic channels, such that the droplet size obtained in these channels is highly controllable. Microfluidic devices also offer versatile means to form complex structures such as non-spherical droplets, anisotropic droplets, or multiple-emulsion “droplets-in-droplets”. These structures can be retained by subsequent droplet solidification, typically achieved through rapid on-chip polymerization, thereby yielding monodisperse particles with complex architecture.

Another application of microfluidic channels encompasses their implementation in scattering experiments. The strategy for this is to place an array of microchannels into an x-ray or neutron beam and to probe the species of interest while flowing through the channels. The channels to be used for this purpose can either have constant dimensions, or they may exhibit varying width and height, allowing the samples to be probed in uni- or biaxially deformed states. As an alternative, a semi-static method aims to use a microfluidic platform which resembles a “parking lot for droplets”, allowing single deformable samples such as droplets or microgel particles to be fixed in addressable positions. Subsequent longterm-observation can serve to monitor ageing or relaxation processes.



Microfluidic fabrication of polymer microgel particles. A flow focusing microfluidic device serves to form monodisperse pre-microparticle droplets, which then serve to template monodisperse microgels.

Research



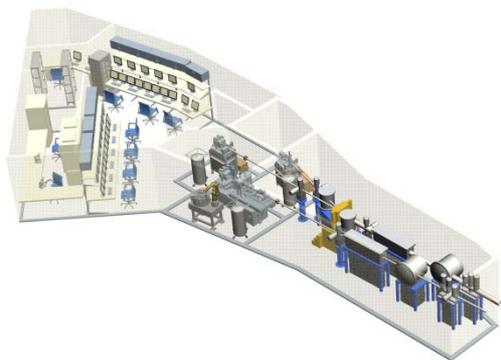
Macromolecular Crystallography (MX)

Uwe Müller, Manfred S. Weiss

Within the past two decades, structural biology has created enormous impact on all biological, biochemical and biomedical research fields. In particular Macromolecular Crystallography (MX) has matured from a few dozen scattered research groups to a field which is now present at every major university and research institution



around the world. This is one of the reasons, why MX beamlines can now be found nowadays at all modern third generation synchrotron sources. The BESSY II based MX research group, which is led by Uwe Mueller and Manfred S. Weiss, operates and develops three MX-beamlines and consists currently of eight researchers and three technical staff members (picture).



The MX activity has been started at BESSY II in strong collaboration with the Free University Berlin in the year 2000. Since 2003 all experimental stations have been used within the regular user operation scheme of BESSY II and have produced more than 500 new protein structures, which are deposited at the Protein structure database PDB (www.rcsb.org/pdb).

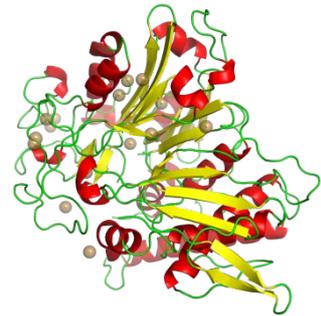
The major task of the MX-group is the provision and the technical as well as scientific support of the experimental resources at a constant high-quality level for the international user community.

The HZB-MX user community currently consists of 43 international research groups, which are running 16% of all photon based research projects at BESSY II. Among these are 12 Berlin based MX-groups. In 2009 the “Joint Berlin MX Laboratory”, a research collaboration between FU-Berlin, the HU-Berlin, the Max Delbrück Centrum Berlin, the Leibniz Institute for Molecular Pharmacology (FMP) and the HZB has been founded to organize the usage, maintenance and development of the MX-beamlines on a collaborative basis.

The BESSY-MX group is also engaged in several research collaborations with regard to the “Joint Berlin MX-Laboratory”, as well as in a number of in-house research project in the field of MX. The two main pillars of these in-house research activities are MX methods development and structure-function relationship of enzymes which are capable of the degradation of environmentally critical soil pollutants.

Phase Determination by Long-Wavelength Sulfur-SAD

In the past years, the use of longer X-ray wavelengths ($\lambda=1.5\text{-}2.5\text{ \AA}$) in MX has gained quite some recognition. The main reason for this is the possibility to determine phases for diffraction data from the sulphur anomalous signal using a method termed sulphur single wavelength anomalous dispersion (S-SAD). To date, about 50 novel protein structures have been determined by the S-SAD method. A particular striking example is the lysosomal 66.3 kDa protein from mouse, since it constitutes the largest protein determined using this method. Furthermore, it crystallizes in a low-symmetry space group (C2), which makes the collection of high-multiplicity data cumbersome and difficult.



Nevertheless, using 1.9 \AA -data collected at the HZB-beamline BL14.2, it was possible to derive the positions of all 22 S-atoms in the 559-residue long protein and use this anomalously scattering S-structure as reference for phase determination. This experiment constitutes a landmark experiment in the area of S-SAD research (figure).

Reference:

- Lakomek, K., Dickmanns, A., Mueller, U., Kollmann, K., Deuschl, F., Berndt, A., Luebke, T. & Ficner, R. (2009). *Acta Cryst.* D65, 220-228.

Group publications (selection):

- K. Gedrich, I.Senkovska, N.Klein, U. Stoeck, A. Henschel, M. R. Lohe, I. A. Baburin, U. Mueller and S. Kaskel. *Eine hochporöse Metall-organische Gerüstverbindung mit zugänglichen Nickelzentren*, *Angewandte Chemie*, 2010, **122**, 8667–8670
- K. S. Paithankar and E. F. Garman. *Know your dose: RADDPOSE*. *Acta Cryst.* **D66**, 381-388
- Faust, S. Pühringer, N. Darowski, S. Panjikar, K. Diederichs, U. Mueller & M. S. Weiss. *Update on the Tutorial for Learning and Teaching Macromolecular Crystallography*. *J. Appl. Cryst.* 2010, **43**, 1230-1237

Listing of all coworkers

- Dr. Martin Bommer
- Dr. Nora Darowski
- Ronald Förster
- Michael Hellmig
- Michael Krug
- Dr. Uwe Mueller
- Dr. Sandra Pühringer
- Michael Steffien
- Dr. Monika Ühlein
- Dr. Manfred S. Weiss

Phase determination Using the Anomalous Signal from Sulfur Atoms

Manfred S. Weiss

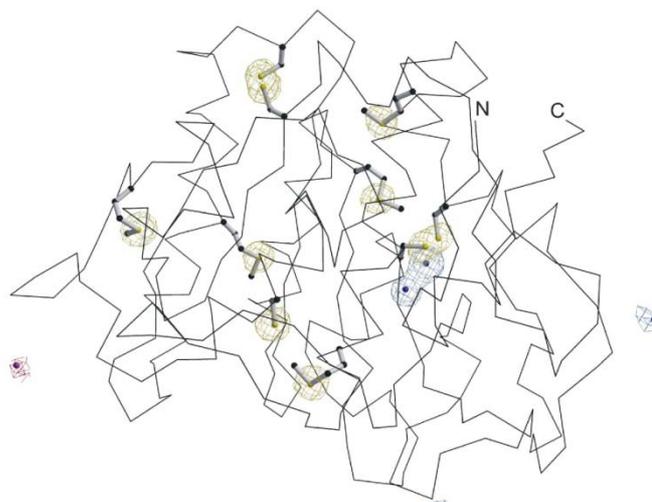
In macromolecular structure determination, the solution of the crystallographic phase problem remains an unsolved issue in the whole process. Over the years, many methods have been developed to derive phases from the diffraction intensities, all of which are grounded upon a modification of the macromolecules under investigation. Since nearly every protein contains sulphur-atoms and every nucleic acid contains phosphorous atoms, it may be anticipated that the anomalous scattering of the S- and P-atoms intrinsically present in the macromolecules, can be used for phase determination. If such an approach would turn out to be commonly applicable, the phase problem in macromolecular crystallography would cease to exist. In the MX-group at the HZB, we are developing tools overcome the major difficulties associated with this **Sulphur Single Wavelength Anomalous Diffraction (S-SAD)** approach.

The main difficulty associated with S-SAD is that the achievable signal is extremely small [1]. The S-SAD related research in the MX-group at the HZB is focussed on two pillars: first, to increase the signal by collecting diffraction data at slightly longer than usual wavelengths (1.8-2.2 Å) and second to measure the intensity differences with a very high precision (about 1%).

A data collection experiment at longer wavelengths is in principle not any different than an experiment at short wavelengths. However, due to the fact that absorption becomes a more and more severe problem as one goes to longer wavelengths, such an experiment has to be carefully planned and carried out. At present, it seems that a data collection wavelength of about 2.0 Å yields the highest anomalous signal-to-noise ratio [2]. With better crystal mounting procedures and better data reduction tools available, however, the hope is that this may shift to even longer wavelengths, where the signal is further increased.

In order to collect diffraction data to the highest possible precision, a number of tools are available at the HZB-MX-beam lines. For instance, BL14.1 is equipped with a kappa-goniometer, which enables the experimenter to orient the crystals and various data collection strategy options help to devise the optimal data collection strategy.

To date, only about 50-100 macromolecular structures have been determined using the S-SAD approach. Two examples are given in references 3 and 4. The current status is that macromolecules crystallized in high-symmetry crystals with a relatively small asymmetric unit and good diffraction properties are amenable to structure determination by S-SAD. We hope, though, that the method can be further developed in order to push the boundaries towards lower symmetries and larger structures.



C_{α} -trace of the enzyme proteinase K. The superimposed anomalous difference electron density clearly shows the positions of all S-atoms (in yellow) of the protein, as well as the bound cations (in blue) and anions (in red).

[1] K. Djinović Carugo *et al.* (2005). *J. Synchr. Rad.* **12**, 410-9.

[2] C. Mueller-Dieckmann *et al.* (2005). *Acta Cryst.* **D61**, 1263-72.

[3] M. S. Weiss *et al.* (2004). *Acta Cryst.* **D60**, 686-95.

[4] K. Lakomek *et al.* (2009). *Acta Cryst.* **D65**, 220-8

Phase Determination Using the UV-Light Induced Radiation Damage

Uwe Mueller

After the collection of an X-ray diffraction data-set from a macromolecule crystal the solution of the so-called “crystallographic phase problem” is the major task, which must be resolved. In order to achieve this, a growing number of methods exist, which we are aiming to extend with the further development of the UV-based radiation induced phasing (UVRIP) method [1]. This experimental technique is focused on specific structural changes of cystine-containing protein crystals, which is due to the irradiation of the specimen with highly-intense UV-radiation (Figure 1). The structural changes can be used to work out a single isomorphous replacement (SIR)-like phasing scheme, which can lead to precise experimental phase information and thus to the access to the three dimensional structure (Figure 2). For this, a native data-set has to be collected before UV-exposure and compared with a second data set collected after the UV-irradiation of the same crystal. At the HZB-MX beamline BL14.1, we have installed all required instruments to carry out such experiments [2,3] and are providing this to the user community. Within this research project, we aim to develop this method to reduce the existing requirements in terms of minimal data-set resolution and to investigate alternative specific damage sites within a macromolecular crystal.

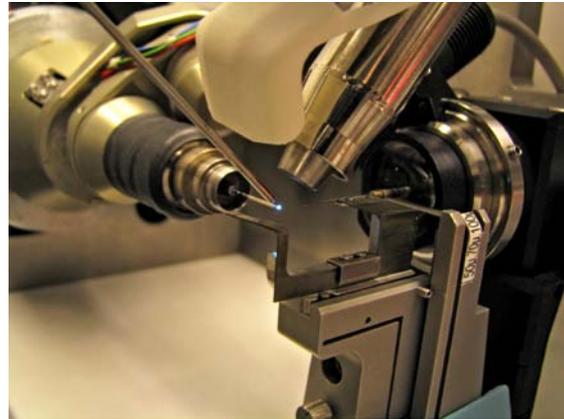


Figure 1: UV-laser setup at BL14.1 during irradiation of an thaumatin crystal

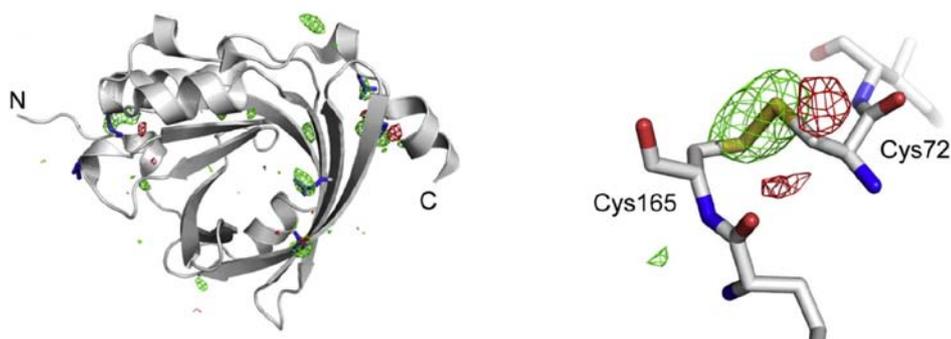


Figure 2: Specifically damaged disulfide-bridges of the protein α_1 -acid glycoprotein

- [1] M. H. Nanao *et al.* (2006). *Structure* **14**, 791-800.
- [2] A. Faust *et al.* (2010). *J. Appl. Cryst.* **43**, 1230-1237.
- [3] D. L. Schoenfeld *et al.* (2008). *J. Mol. Biol.* **384**, 393-405.

Diffraction-Based Screening to Rapidly Characterize Biological Crystals in their Native Environment

Karthik S. Paithankar

One of the challenges in macromolecular crystallography is the identification of suitable crystals for diffraction data collection. In the past decade structural genomics projects and pharmaceutical companies have successfully created pipelines to produce and purify proteins in a rapid manner. Crystals from these proteins are grown in crystallization trays being able to harbour 96, 192, or 288 different experimental conditions, using automated liquid handling systems (Figure 1). So far crystals are pre-selected by optical imaging but this method does not provide any information about diffraction properties of the specimen. Using a microscope it is often not possible to distinguish salt and other small molecule from protein crystals. In addition, the first crystals obtained in these experiments are usually very small and fragile.

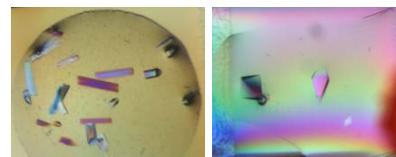


Figure. 1 Crystals of the proteins insulin and lysozyme



Figure 2 A crystallization plate with samples ready for X-ray analysis

One alternative to the classical imaging method is the rapid identification of crystallization conditions that provide diffraction quality crystals *in situ* at room temperature by X-ray diffraction [1].

Within our group, we are developing and applying such methods by exposing the crystallization trays directly to X-ray beam at room temperature. using a robot.

The hardware implementation consists of a 6-axis industrial robot arm, which can handle crystallization plates (Figure 1). The crystal, which is grown inside the drop in the plate, can thus be directly exposed by the X-ray beam (Figure 3). Only few X-ray images are required to gain information about the crystal characteristics, such as the unit cell, symmetry and mosaicity.

Thus, instead of undergoing the long and exhaustive process of growing hundreds of crystals and testing each crystal in the X-ray beam, a single crystallization tray can be exposed to find the diffraction characteristics of all freshly grown crystals at ones.

Note in addition: Recently, a joint BMBF-funded research project has been initialized between the HZB and the Russian Academy of Sciences and the Freie Universität Berlin to develop this method.

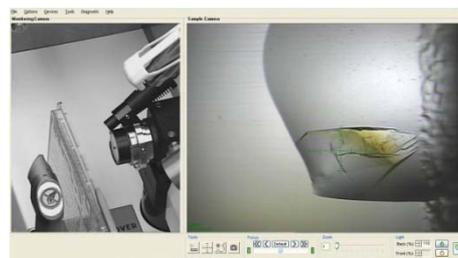


Figure 3 A view of the crystal after X-ray irradiation

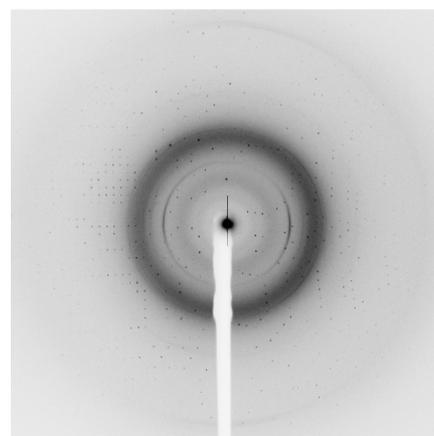


Figure 4 Diffraction image from a protein crystal sitting in a plate

[1] Jacquamet *et al.* (2009). *J. Synchr. Rad.* **16**, 14-21.

Structure and Function of Protein Complexes from the Spliceosome

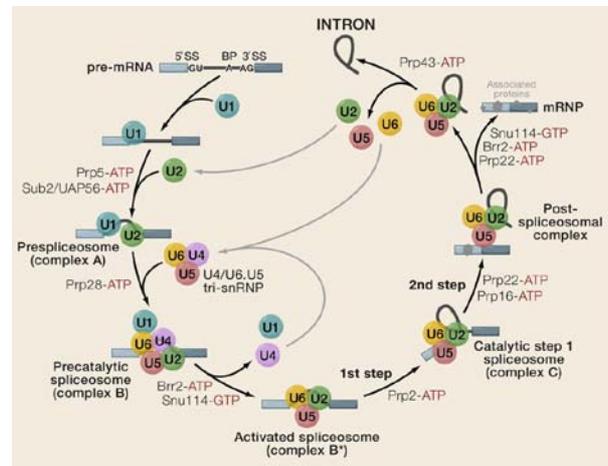
Sandra Pühringer

The spliceosome is a large and highly dynamic RNA-protein machine that deletes unimportant (so called “non-coding”) regions from the DNA – a process called splicing. For each splicing event, a spliceosome is assembled *de novo* on the pre-mRNA, extensively remodeled and, after the deletion of the non-coding regions, disassembled in an ordered fashion (Figure).

As in all complex cellular processes splicing is also known to be a potential source of errors during gene expression. Several diseases are known that are directly linked to errors during pre-mRNA splicing (e.g. retinitis pigmentosa) and even specific forms of cancer are potentially caused by spliceosomal anomalies.

In humans, about 200 proteins, five small nuclear RNAs and the pre-mRNA intimately participate in the splicing process. During spliceosome assembly, catalysis and disassembly, many protein-protein, protein-RNA and RNA-RNA interactions are formed and broken in a controlled manner [1]. Presently, the exact sequence of remodeling steps and the functional roles of individual components of the spliceosome during assembly and catalysis are not fully understood.

In close collaboration with member institutes of the Joint Berlin MX-Laboratory, we screen for low molecular weight substances, which hyper-stabilize crucial protein-protein interactions that normally form only transiently during splicing. In a radically new approach we are aiming towards the identification of compounds, which bind at the interface of protein complexes and have the potential to hyperstabilize these interfaces. Such compounds would help in biochemical studies of spliceosomal assemblies since they arrest the molecular machine at a certain stage and they would also provide the basis for new therapeutic approaches. It must be anticipated that for such a project a very large number of diffraction data sets must be collected in order to identify suitable compounds. Therefore, the environment created by and the possibilities within the Joint Berlin MX-laboratory are ideally suited to carry out this project.

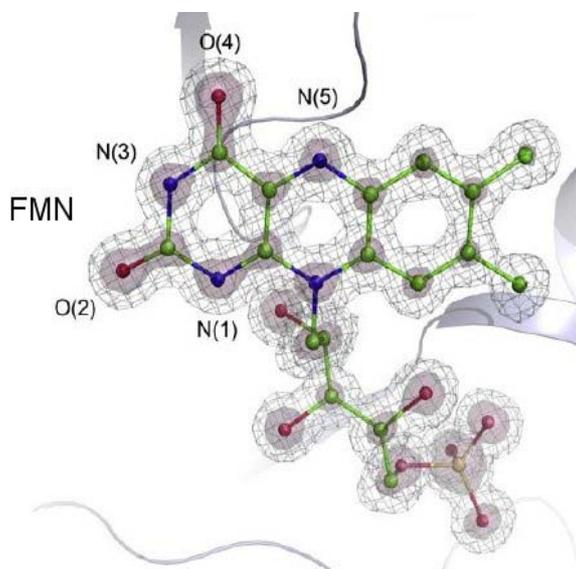


Schematic representation of the spliceosomal assembly. Adapted from [1].

[1] M. C. Wahl *et al.* (2009). *Cell*. 136, :701-718.

Xenobiotic Reductase A of *Pseudomonas Putida*

Michael Krug



The FMN molecule in the active site of XenA and its corresponding electron density. Oxygen atoms are shown in red, nitrogen atoms in blue, carbon atoms in green and phosphorus in yellow

Xenobiotics are substances that can be found in biological systems but are not supposed to be there (xenos=foreign, bios=life). Due to continuous evolutionary pressure, microbes have developed a diversity of pathways to cope with and to degrade xenobiotics such as aromatic and hetero-aromatic compounds.

Pseudomonas putida 86 was isolated near a coal tar factory (Rütgerswerke, Castrop-Rauxel) in Germany. Xenobiotic reductase A (XenA) of *P. putida* 86 is involved in the degradation of quinoline, an ubiquitous soluble, heteroaromatic

pollutant with cancerogenic properties. XenA contains flavin mononucleotide (FMN) resulting in a yellowish color of the enzyme. FMN is produced from vitamin B₂ in the cell and plays an important role in reactions comprising electron transfers.

XenA catalyzes the reduction of various substrates. The reaction of XenA can be divided into two half-reactions. In the first half-reaction, the oxidized enzyme is reduced by NADH, a reducing agent of living cells. In the second half-reaction, the enzyme itself can reduce different substrates whereby it gets oxidized. Atomic resolution crystal structures of XenA showed that the redox-sensitive bond lengths of the FMN are neither typical for oxidized nor for reduced flavins but are in between the distances expected for either one of the oxidation states. It is therefore likely that the synchrotron radiation reduced at least parts of the protein molecules during data collection leading to a mixture of oxidized and reduced molecules in the crystal.

In order to get a better understanding of how the enzyme works, it is essential to obtain insights into the structural differences between the oxidized and the reduced state of XenA. Using adapted data collection and data merging strategies, we are attempting to separate the data representing the oxidized state of the enzyme from those representing the reduced state. Consequently, we will obtain high-resolution structural information for both oxidation states.

[1] O. Spiegelhauer *et al.* (2010). *J. Mol. Biol.* **396**, 66-82.

[2] J. Griese *et al.* (2006). *J. Mol. Biol.* **361**, 140-152.

[3] F. Spiegelhauer *et al.* (2009). *Biochemistry* **48**, 11412-11420.

Biological Degradation of Halogenated Organic Pollutants: Enzymes, Structures & Mechanisms

Martin Bommer, Holger Dobbek (Humboldt Universität zu Berlin)

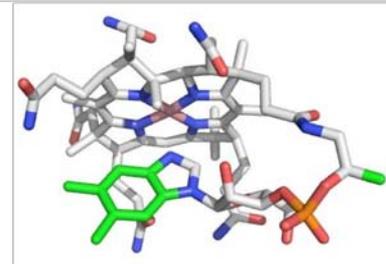
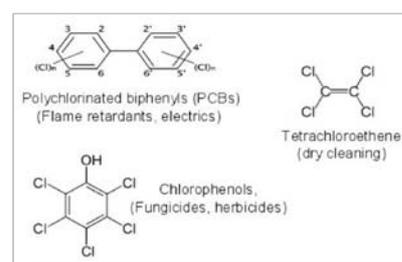
The largest group of priority “persistent organic pollutants” identified by the United Nations Environment Program is the one comprising the class of halogenated, and in particular chlorinated, organic compounds. These are produced globally at industrial scale and are persistent in the soil at contaminated industrial sites. In addition, a number of undesirable compounds are also produced biologically. Chloromethane is mainly produced naturally and is a major contributor to ozone layer depletion. Similarly, most of the above-mentioned industrial pollutants also exist in the natural environment, albeit in much lower quantities. Conversely, the world's finely balanced ecosystem also provides a degradation route for these chemicals.

Within the BESSY-MX team, we are studying the dehalogenation reactions, which yield the harmless halide salt and non-chlorinated hydrocarbon. In nature, these are one step in complex enzymatic reaction cascades that provide energy for soil bacterial (dehalorespiration).

We are looking at the single enzyme (the biological catalyst) and its reaction mechanisms at atomic scale. Two of these are reductive dehalogenase, which dechlorinates the chemicals shown on the right, and chloromethane dehalogenase. Both enzymes use vitamin B12 (bottom) as co-factor, a complex corrinoid ring with a central redox-active cobalt ion (pink in the structure).

We hope to complete this picture and include the surrounding amino acid scaffold provided by the enzyme, which shields the activated cobalt and gives this reaction its specificity.

This will be investigated by X-ray diffraction using the synchrotron radiation at BESSY II and by time-resolved biochemical characterization at the Laboratory of Prof. Holger Dobbek at the Humboldt Universität zu Berlin.



Top: The Buna Chemical works at the Saale River in 1990, which provide a model system for dehalogenation
Centre: selected organic pollutants and their original applications
Bottom: An atomic model of Vitamin B12, which forms the core of the dehalogenase enzymes

[1] Hollinger *et al.* (1999) *FEMS Reviews* **22**, 383-398

[2] Futagami *et al.* (2008). *The Chemical Record* **8**, 1-12.

Biophysics

Thomas Hauß

The biophysics group provides expertise on the investigation of proteins and especially biological membranes with neutron scattering techniques to study the structure and dynamics of these systems. The research topics include the investigation of biological (model) membranes with embedded peptides and proteins and proteins adsorbed to colloidal nano-particles.

Our group runs the neutron diffractometer V1 (Membrane Diffractometer) and the biophysics user laboratory (BioLab) on the Lise-Meitner-Campus Wannsee. The BioLab provides state-of-the-art instrumentation and techniques for guest scientists preparing and complementing their neutron scattering experiments on-site with a variety of biophysical and biochemical methods. One aspect of our work is to develop and provide unique and specialised sample environment, one example is a dedicated excess water cell to study membrane structures near physiological conditions. The Membrane Diffractometer served 10 external user experiments in 2010, 4 in-house experiments and 2 investigations with our long-term cooperation partner N.A. Dencher from the Technische Universität Darmstadt.

The main topics of the in-house research of our biophysics group are membrane biophysics and protein dynamics. With our longstanding expertise in this field, we are developing new techniques to study these systems by neutron diffraction, reflectometry and small angle scattering (both neutron and X-ray) very close to physiological conditions. The use of different scattering methods highlights specific characteristics of the systems under investigation.

The emerging trend for the explanation of neuro-degeneration in Alzheimer's disease imputes the cause of neurotoxicity to the interaction of soluble forms of amyloid- β peptide ($A\beta$) with neural cells and cell membranes. In a series of neutron diffraction and neutron small angle experiments we established that the toxic fragment $A\beta(25-35)$ is able to penetrate and perturb lipid membranes and that the $A\beta$ peptide induces membrane fusion. With quasi-elastic neutron scattering we observed an accelerated lateral diffusion of lipids in membranes doped with $A\beta(25-35)$.

Very recently we developed in cooperation with J. Pieper (University of Tartu, Estonia) a new method to study in situ time resolved protein dynamics with a novel laser-pump – neutron-probe experiment. Our findings showed for the first time a modulation of the protein dynamics during a working cycle of a protein, here bacteriorhodopsin. Our goal is to characterise this modulation in dependence of important environmental parameters like hydration, temperature, pH, lipid composition, and others. To adopt this new method to proteins, which are not directly activated by a photon, the use of caged compounds, such as caged Ca, H^+ , or ATP in a flow-cell will be investigated.



Membrane Diffractometer V1

Sample mounted on V1

Excess water cell

Another area of interest in the biophysics group is the understanding of the mechanism of protein adsorption onto nanoparticles, as elucidated by their thermodynamic parameters. Protein adsorption onto nanomaterials is of interest in diverse applications including nanomedicines, food and waste processing, water purification and diagnostics. Isothermal titration calorimetry is used to determine the protein binding isotherm with varying temperature, salt and pH. The resulting changes in entropy, enthalpy and equilibrium constant can provide information about the relative importance of electrostatic and hydrophobic contributions to adsorption.

List of co-workers:

Dr. Thomas Hauß
Alisa Becker, PhD
Dr. Alexei Plotnikov
Nicole Welsch
Elzbieta Charkiewicz
Dennis Schmidt
Björn Drobot
Alexandra Graebert
Luigi Sparacio

Selected publications 2009-2011:

- 1) Pieper J, Buchsteiner A, Dencher NA, Lechner RE, Hauß T, *Light-induced modulation of protein dynamics during the photocycle of bacteriorhodopsin*. Photochem Photobiol 2009, **85**, 590-597.
- 2) Schröter A, Kessner D, Kiselev MA, Hauß T, Dante S, Neubert RHH: *Basic nanostructure of stratum corneum lipid matrices based on ceramides [EOS] and [AP]: a neutron diffraction study*. Biophys J 2009, **97**, 1104-1114.
- 3) Welsch N, Wittemann A, Ballauff M: *Enhanced activity of enzymes immobilized in thermoresponsive core-shell microgels*. J Phys Chem B 2009, **113**, 16039-16045.
- 4) Buchsteiner A, Hauß T, Dante S, Dencher NA: *Alzheimer's disease amyloid-beta peptide analogue alters the ps-dynamics of phospholipid membranes*. Biochim Biophys Acta 2010, **1798**, 1969-1976.
- 5) Ryabova NY, Kiselev MA, Dante S, Hauß T, Balagurov AM: *Investigation of stratum corneum lipid model membranes with free fatty acid composition by neutron diffraction*. Eur Biophys J 2010, **39**, 1167-1176.
- 6) Welsch N, Ballauff M, Lu Y: *Microgels as Nanoreactors. Applications in Catalysis* 2010, **234**, 129-163.
- 7) Graebert A, Schmidt D, Kyriakopoulos A: *Selenoproteome of the nuclear envelope of JTC-15 cells*. International Journal of Trends in Medicine 2011: *accepted*.
- 8) Wagner CS, Shehata S, Henzler K, Yuan J, Wittemann A: *Towards nanoscale composite particles of dual complexity*. Journal of Colloid and Interface Science 2011, **355**, 115-123.

Time Resolved Protein Dynamics

Thomas Hauß, J. Pieper*, R. Lechner, A. Buchsteiner[†], N.A. Dencher[‡]

Proper functioning of a protein requires a well-defined three-dimensional structure. However, a protein is not a static entity; the working protein often undergoes larger conformational changes within microseconds and usually it needs a certain internal flexibility, which is provided by stochastic structural fluctuations on the picoseconds time scale. These fluctuations are represented by anharmonic vibrational and local diffusive motions of small molecular subgroups in the bulk of the protein as well as on its surface. For the understanding of the structure-function relationship on an atomic level it is of great importance to increase our knowledge about these dynamical properties. Inelastic and quasi-elastic neutron scattering techniques, in the past, contributed much to the present knowledge, however, the correlation between internal protein dynamics and functionality has only been studied indirectly in steady-state experiments by variation of external parameters like temperature or hydration.

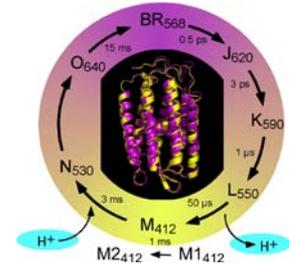


Fig. 1 Photocycle of Bacteriorhodopsin and its structure in ground-state (purple) and M-intermediate.

We have developed in cooperation with J. Pieper (University of Tartu, Estonia) a novel laser-pump – neutron-probe experiment, which combines in-situ

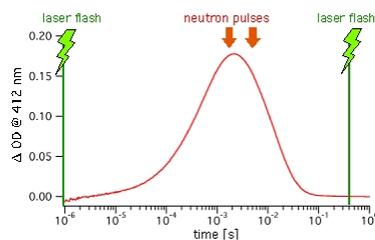


Fig.2 Time course of the laser-pump – neutron-probe experiment: the photocycle is triggered by a laser flash and monitored with the raise and decay of the M-intermediate with optical spectroscopy at 412 nm. The neutron probe pulses test the ps-dynamics at ~1 ms where BR protein is in its M-intermediate.

optical activation of the function of a membrane protein, here bacteriorhodopsin, with a time-dependent sampling of the modulation of its protein dynamics using quasi-elastic neutron scattering. Our results demonstrate for the first time temporary alterations in the protein dynamics after triggering the working cycle of the membrane protein. This observation is a direct proof for the functional significance of protein structural flexibility, in connection with the large-scale conformational changes in the protein structure occurring during the operation of a “molecular machine”. Functionally important modulations of protein dynamics as observed here are surely of relevance for most of the proteins that exhibit conformational changes in the time course of its functional operation. Therefore, our next goal is to adapt our newly developed laser-pump – neutron-probe technique to proteins which can indirectly be triggered by caged compounds, of caged calcium.

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- [2] Pieper J, Buchsteiner A, Dencher NA, Lechner RE, Hauß T, *Light-induced modulation of protein dynamics during the photocycle of bacteriorhodopsin*. Photochem Photobiol 2009, **85**,590-597.
- [3] Seelert H, Dani DN, Dante S, Hauß T, Krause F, Schäfer E, Frenzel M, Poetsch A, Rexroth S, Schwaßmann HJ *et al*, *From protons to OXPHOS supercomplexes and Alzheimer's disease: Structure-dynamics-function relationships of energy-transducing membranes*. Biochim Biophys Acta 2009, **1787**, 657-671.

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Interactions between Proteins and Colloidal Particles

Alisa Becker, Nicole Welsch

The interface of materials science and biology is emerging as a major research focus, in particular, nanomaterials in medicine and biotechnology are expected to address many medical and biological problems. In these complex environments, nanoparticles come into contact with proteins, which can be adsorbed onto the particle surface. This changes the surface chemistry of the particle, and can also change the activity and conformation of the protein.

We investigate protein adsorption onto nanomaterials using two types of model nanoparticles. Spherical polyelectrolyte brushes (SPB) synthesized at the HZB consist of a solid polystyrene (PS) core onto which monodisperse polyelectrolyte chains are attached. Therefore, the adsorption of proteins on SPBs is electrostatically controlled. The second type of particles are core-shell microgels based on poly(N-isopropylacrylamide) (PNiPA). Herein, the PS cores are surrounded by a crosslinked PNiPA shell, which can be copolymerized with charged comonomers to produce microgels of different charged states. PNiPA exhibits a lower critical solution temperature (LCST) close to body temperature, where the microgel network shrinks and swells upon temperature changes (Fig. 1). Therefore, these smart core-shell microgels have the potential to adsorb and release proteins in a controlled way.

Calorimetric and scattering methods (e.g. Isothermal Titration Calorimetry, ITC and Small Angle X-ray

Scattering, SAXS) are used to characterize the thermodynamics of the spontaneous adsorption process, as well as the spatial distribution of the proteins within the SPBs and microgels (Fig. 2).

In addition, kinetic experiments with immobilized enzymes are performed to determine the impact of immobilization on the enzymatic activity (Fig. 3). In all cases investigated the activity of adsorbed enzymes is retained or even increased upon adsorption. Furthermore, temperature-dependent kinetic experiments indicate that the catalytic activity of enzymes immobilized in thermosensitive microgels can be altered by the volume phase transition of the smart carriers.

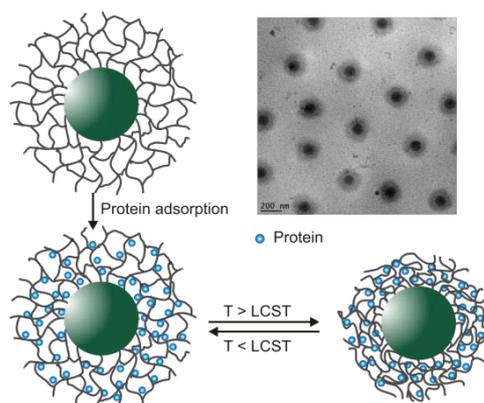


Figure 1: Top left and bottom: Schematic representation of the adsorption of proteins on PS-PNiPA microgel particles. Top right: Cryo-TEM image of these particles in the swollen state.

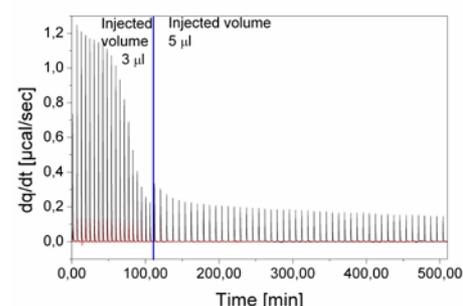


Figure 2: ITC data for titration of lysozyme into a solution of negatively charged microgels at pH 7.2 and at 25°C.

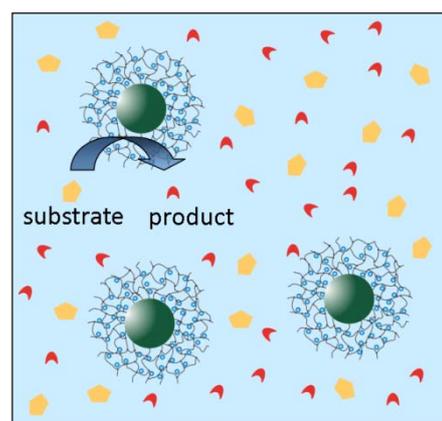


Figure 3: Schematic representation of the catalysis with immobilized enzymes.

[1] Henzler, K. *et al*, Phys. Rev. Lett., 2008, **100**, 158301

[2] Henzler, K. *et al*, J. Am. Chem. Soc., 2010, **132**, 3159-3163

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Dynamics of Phospholipid Membranes Influenced by the Alzheimer's Disease Amyloid- β Peptide Analogue

Thomas Hauß, Alexandra Buchsteiner*

We investigate the influence of the neurotoxic Alzheimer's disease peptide amyloid- β and various fragments of it on the dynamics of phospholipid membranes by means of quasi-elastic neutron scattering (QENS) in the picosecond time-scale.

Samples of pure phospholipids (DMPC/DMPS) and samples with amyloid- β (25-35) peptide included have already been compared. With two different orientations of the samples the directional dependence of the dynamics was probed. The sample temperature was varied between 290 K and 320 K to cover both the gel phase and the liquid-crystalline phase of the lipid membranes.

The model for describing the dynamics combines a long-range translational diffusion of the lipid molecules and a spatially restricted diffusive motion.

Amyloid- β (25-35) peptide affects significantly the ps-dynamics of oriented lipid membranes in different ways. It increases the lateral diffusion velocity especially in the liquid-crystalline phase. This is very important for all kinds of protein-protein interactions which are enabled and strongly influenced by the lateral diffusion such as signal and energy transducing cascades. Amyloid- β (25-35) peptide also increases the local lipid mobility as probed by variations of the vibrational motions with a larger effect in the out-of-plane direction.

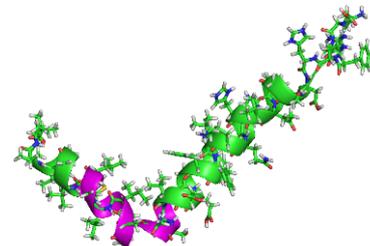
Thus, the insertion of amyloid- β (25-35) peptide changes not only the structure of phospholipid membranes as previously demonstrated by us employing neutron diffraction but also the dynamics inside the membranes.

The amyloid- β (25-35) peptide induced membrane alteration even at only 3 mol% might be involved in the pathology of Alzheimer's disease as well as be a clue in early diagnosis and therapy.

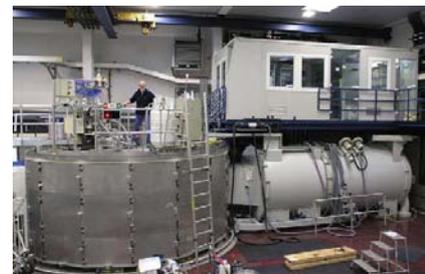
In future, we plan to investigate longer fragments of amyloid- β and its influence on natural membranes rather than phospholipid model membranes.

[1] A. Buchsteiner, T. Hauß, S. Dante, N.A. Dencher: *Alzheimer's disease amyloid- β peptide analogue alters the ps-dynamics of phospholipid membranes*, Biochimica et Biophysica Acta 2010, **1798**, 1969-1976

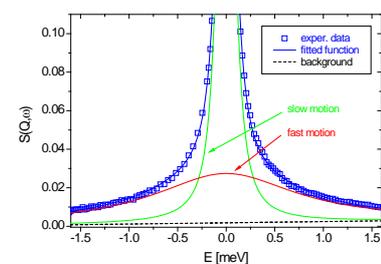
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Heinrich-Damerow-Str. 4 D-06120 Halle Germany



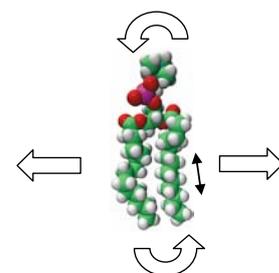
structure of amyloid- β (1-42) in aqueous solution (PDB ID: 1IYT) with the neurotoxic fragment amyloid- β (25-35) highlighted



time-of-flight spectrometer NEAT



typical QENS spectrum with two different motions



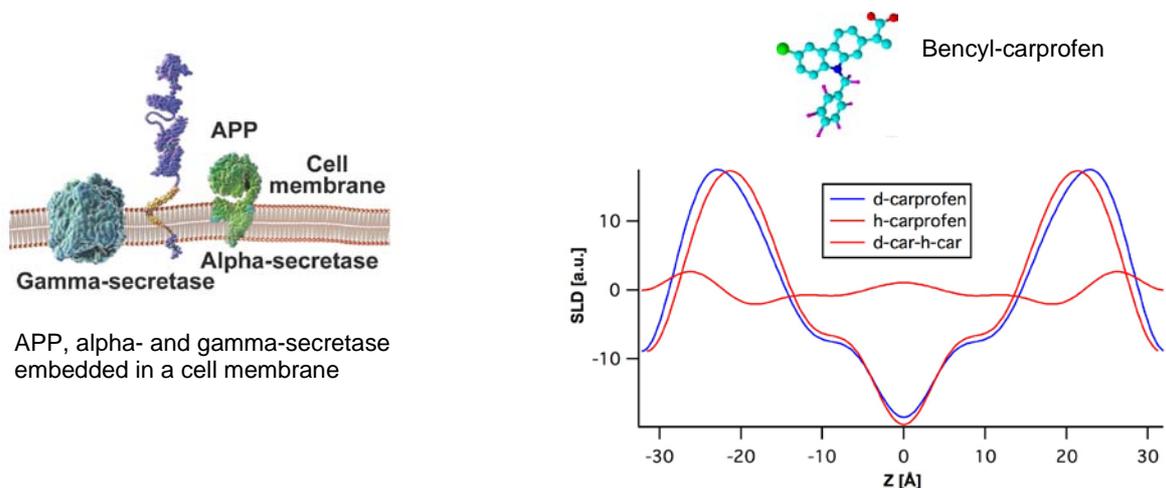
example of a phospholipid molecule with possible modes of motion indicated

Interaction of a γ -Secretase Modulator with Model Lipid Membranes

Thomas Hauß¹, Boris Schmidt², Norbert A. Dencher³

Recently, new strategies were developed to find a therapeutic approach to Alzheimer's disease [1,2,3]. The targets of the new approaches are enzymes responsible for the cleavage or modification of the amyloid precursor protein (APP) into the neurotoxic peptides β -amyloid with 40 to 43 amino acids. In the group of Professor Schmidt, TU-Darmstadt newly designed inhibitors and modulators for β - and γ -secretase were successfully tested in cultured cells and in mice [1,2]. The cleavage site of γ -secretase is in the hydrophobic core of the cell membrane, for that reason the inhibitor of γ -secretase is lipophilic.

We investigated by neutron diffraction the interaction of a newly synthesised and specifically deuterated γ -secretase modulator, a carprofen derivative, with lipid membranes. We established a suitable protocol for the preparation of biological highly relevant membrane models consisting of POPC, sphingomyelin, and cholesterol. This lipid mixture exhibits a change in lattice spacing, indicating a phase transition, at temperatures between 10°C and 40°C, with and without the inhibitor. The neutron diffraction experiments revealed the localization of the deuterated inhibitor in the membrane lipids as difference in the scattering length density profiles. The difference is calculated from samples of membrane lipids mixed with the protonated or selectively deuterated inhibitor carprofen, respectively, at two different contrast points (8% D₂O, 20% D₂O in the aqueous atmosphere) at 15°C. The maxima in the difference density profile at $z = \pm 2.5$ nm are attributed to the location of the deuterated label. The tentative interpretation is, that the modulator with its deuterated benzyl ring resides in the head-group region of the lipid membrane close to the phosphate group of the lipids.



Neutron scattering length density profiles of lipid membranes with deuterated and protonated carprofen derivatives, respectively, and its difference indicating the position of the deuterated benzyl ring

[1] Rajendran et al., Efficient inhibition of the Alzheimer's Disease β -secretase by Membrane targeting, *Science* **320**, 520 (2008)

[2] Kukar et al., Substrate-targeting γ -secretase modulators, *Nature* **453**, 925 (2008)

[3] Schilling et al., Glutaminy cyclase inhibition attenuates pyroglutamate A and Alzheimer's disease-like pathology, *Nature Medicine* **14**, 1106 (2008)

- 1) Helmholtz-Zentrum Berlin, Institut für weiche Materie und funktionale Materialien
- 2) Technische Universität Darmstadt, Clemens Schöpf-Institut, Organische Chemie,
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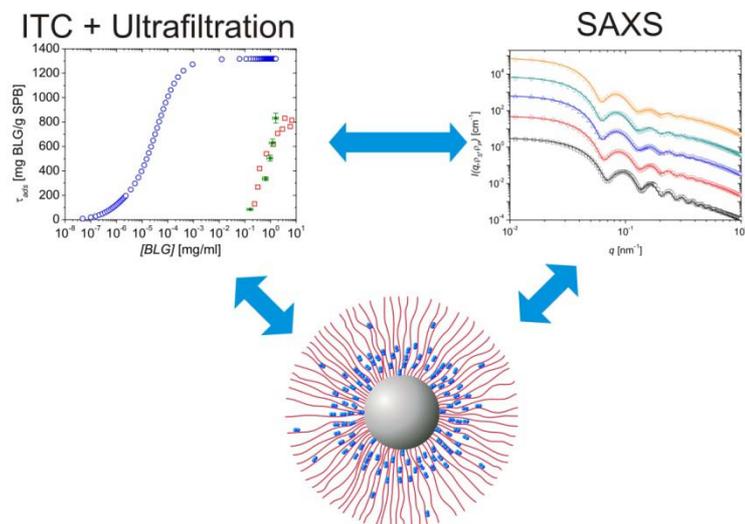
Interaction Strength between Proteins and Polyelectrolyte Brushes: A Small Angle X-Ray Scattering Study

K. Henzler^{1,2}, B. Haupt^{1,2}, S. Rosenfeldt², L. Harnau³, T. Narayanan⁴, M. Ballauff^{1,4}

The interaction of proteins with different surfaces is of central interest for biotechnology and medical science. Here, we present an investigation of the adsorption of β -lactoglobulin (BLG) onto spherical polyelectrolyte brushes (SPB) by small angle X-ray scattering, isothermal titration calorimetry and extensive ultrafiltration.

This demonstrates for the first time that an in-depth understanding

of the interaction strength of proteins with polyelectrolyte brushes can be obtained by the proper combination of different analytical methods. The amount and distribution of the protein insight the brush layer can be determined by small angle X-ray scattering (SAXS).[1] Furthermore, the SAXS measurement shows that a certain amount of the protein molecules form linear aggregates in the adsorbed state of about six monomer units. On the other hand, isothermal titration calorimetry (ITC) provides the opportunity to investigate the thermodynamics of the protein adsorption as well as the amount of adsorbed protein at the equilibrium.[2] The amount of adsorbed BLG determined by ITC and SAXS can be compared. The third method used in this investigation is the extensive ultrafiltration.[3] By this method the amount of tightly bound BLG can be determined. From the SAXS analysis the amount of adsorbed protein in different parts of the polyelectrolyte layer can be calculated. These results can be compared to the data obtained by ultrafiltration. It is found that the proteins which are bound in the outer part of the brush layer can be washed out by ultrafiltration.



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[2] Henzler et al., Adsorption of beta-Lactoglobulin on Spherical Polyelectrolyte Brushes: Direct Proof of Counterion Release by Isothermal Titration Calorimetry, *J. Am. Chem. Soc.* **132**, 3159 (2010)

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- 1) Helmholtz-Zentrum Berlin, Institut für weiche Materie und funktionale Materialien
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- 4) European Synchrotron Radiation Facility Grenoble

Colloid Physics

Guenter Goerigk, Daniel Clemens

The colloid physics group is providing instrumentation for investigations of structure and dynamics of large scale structures using neutrons, X-rays and light as suitable probes. We continue to develop our techniques in order to access important outer parameters as temperature, shear forces, etc. that have an influence on the systems under investigation. Our instrumentation includes small-angle scattering (V16/VSANS) and spin-echo spectroscopy (V5/SPAN) on the neutron side, we participate in the operation of an anomalous small-angle X-ray scattering (ASAXS) beam line at the BESSY II synchrotron that belongs to the institute F-11.

The colloid physics group is also in charge of the colloid lab. Here we are running additional methods as e.g. static and dynamic light scattering, zeta-sizer as well as rheometers that supplement investigations by the beamlines of our users. These facilities are open to all users of the beamlines of F-12. Additional techniques are assembled in the newly founded **Joint Laboratory of Structural Research (JLSR)** which is run in close collaboration with the Institute of Physics of the Humboldt University. A new cryogenic transmission electron microscope (cryo-TEM) will be installed there in April 2011. This instrument allows us to analyze suspensions of colloidal and biological systems in aqueous environment. The results obtained in this way can directly be compared to similar studied done with the X-ray microscope (see the description of the X-ray microscopy group). Moreover, the analysis done in real space is compared to small-angle experiments done by SANS and SAXS.

The group is engaged in selected research activities, e.g. in the field of applied small-angle scattering of colloids: microdomains in lipid membranes, critical phenomena at mesoscopic scale, comparison to theoretical predictions for polyelectrolyte brushes and the rheology of hard sphere suspensions. Moreover, we have an intense cooperation with research groups in the field of solar cells, plasmonics and catalysis within and outside HZB.

Small-Angle Scattering with Synchrotron Radiation and Neutrons – Precise Experimental Techniques for the Analysis of Fluctuations and Critical Phenomena

Small-Angle Scattering (SAS) experiments average over a large sample volume and give structural and quantitative information of high statistical significance on a mesoscopic length scale between 1 and hundreds of nanometers, which can be correlated with macroscopic physical and chemical parameters of the analyzed materials. The materials under investigation cover a wide range of different scientific fields for instance, macromolecules in solution, suspensions, metal nanoparticles, composites, membranes, alloys, semiconductors, glasses...).

Synchrotron radiation and neutrons provide extraordinary powerful tools for SAS experiments. By use of Anomalous Small-Angle X-ray Scattering (ASAXS) synchrotron radiation employs the energy tunability in the vicinity of the K- and LIII-absorption edges of most of the elements giving access to the element-specific structural and quantitative characterization of the samples under investigation. From a series of publications of the last years it has been shown, that tremendous quantitative information about chemical concentrations in highly diluted chemical solutions can be obtained by q-ASAXS when employing the so-called Resonant Invariant (RI). From the integral (RI) in Figure 1 the amount of Sr counter ions localized in collapsed subdomains of polyacrylate with respect to the total amount of Sr-cations in the solvent was deduced, while approaching the systems phase boundary.

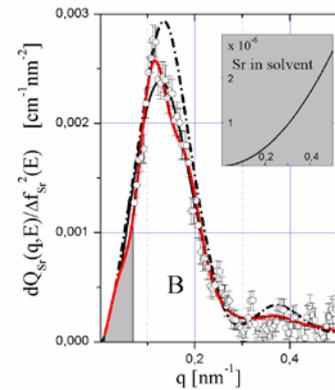


Fig. 1 Resonant Invariant of Sr counter ions localized in collapsed subdomains of polyacrylates.

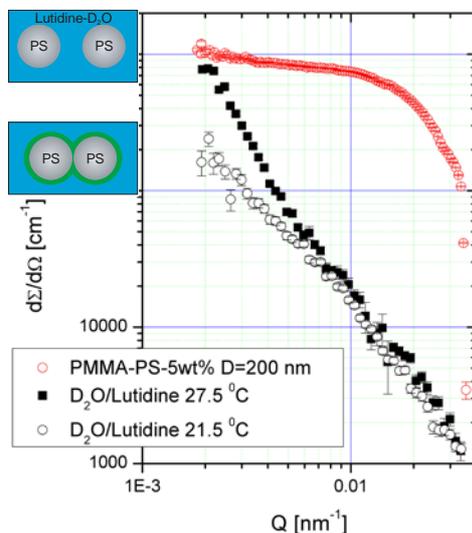


Fig.2: 200 nm colloids at different temperatures near the critical point with respectively without critical Casimir forces

The influence of critical fluctuations becomes visible at 27.5 °C.

A point of special interest is the combination of cryo-TEM with scattering methods. Recently, we analyzed nano-crystals from polyethylene in collaboration with the group of Prof. S. Mecking, University of Konstanz. The shape and the size distribution of the particles was assessed directly by cryo-TEM while their internal structure was analyzed by SAXS including contrast variation. In this way a full structural model of these particles that present the smallest polymer crystals ever synthesized could be achieved. Moreover, the lamellar thickening of the nanocrystals after thermal annealing could be analyzed.

Beside other applications Small Angle Neutron Scattering provides contrast variation for soft matter research by deuteration or H₂O/D₂O mixtures and - by use of special neutron optics – the analysis of large scale structures up to the micro meter length scale by so-called Very Small-Angle Neutron Scattering (V-SANS). Figure 2 summarizes first results from basic research upon the Critical Casimir Effect of large colloids in critical Lutidin/D₂O mixtures. The red scattering curve represents PS-colloids of 200 nm diameter obtained from the JCMS-instrument KWS-3 at the Research reactor FRM II of Technische Universität München. The black and the blue symbols show the scattering curves of the critical mixture at different

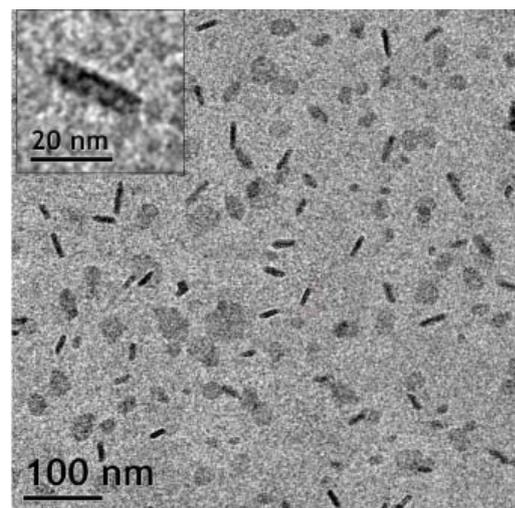


Fig. 3 Cryo-TEM micrograph of an aqueous suspension of polyethylene nanocrystals. The inset displays an enlarged picture of one crystal. This analysis has been combined with SAXS to arrive at a full model of the smallest polymer crystals made so far. Taken from ref. [14] selected publications.

Coworkers:

Dr. Beate-Annette Brüning

Dr. Daniel Clemens

Dr. Günter Goerigk

Dr. Sylvain Prévost

Christian Rabe

Christian Schneider

Miriam Siebenbürger

Dr. Ralf Stehle

Dr. Karsten Vogtt

**Selected Publications in 2009-2010:**

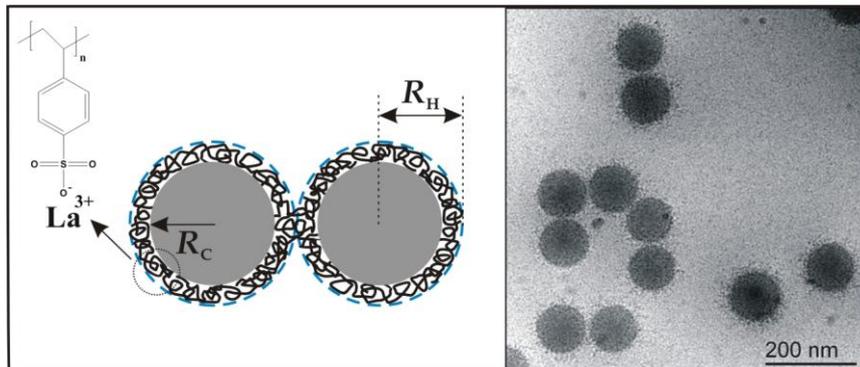
1. Hoffmann, M.; Jusufi, A.; Schneider, C.; Ballauff, M. *J. Colloid Interface Sci.*, **2009**, 338(2), 566-572: "Surface potential of spherical polyelectrolyte brushes in the presence of trivalent counterions"
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3. S. Lyonnard, Q. Berrod, B. Brüning, G. Gebel, A. Guillermo, H. Ftouni, J. Ollivier, B. Frick; Perfluorinated surfactants as model charged systems for understanding the effect of confinement on proton transport and water mobility in fuel cell membranes. A study by QENS; *Eur. Phys. J ST* 189: 205-216 (2010).
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8. M. Hoffmann, M. Siebenbürger, L. Harnau, M. Hund, C. Hanske, Y. Lu, C.S. Wagner, M. Drechsler, M. Ballauff, "Thermoresponsive colloidal molecules." *Soft Matter*, **6**, 1125-1128 (2010).
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11. Vogtt, K., Jeworrek, C., Garamus, V. and Winter, R. (2010), *Microdomain Formation in Lipid Vesicles: Structure and Distribution Assessed by Small Angle Neutron Scattering*, *Journal of Physical Chemistry B*, **114(16)**, 5643–5648 (2010).
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13. Goerigk, G., and Varga, Z., *J. Appl. Cryst.* 2011; **44** doi:10.1107/S0021889811000628
14. C. N. Rochette, S. Rosenfeldt, K. Henzler, F. Polzer, M. Ballauff, Q. Tong, S. Mecking, M. Drechsler, T. Narayanan, L. Harnau; Annealing of single lamella nanoparticles of polyethylene, *Macromolecules*, submitted

The Colloidal Stability of Spherical Polyelectrolyte Brushes

Christian Schneider

Polyelectrolyte brushes are systems in which long polyelectrolyte chains are densely attached to planar or curved surfaces. Attaching long polyelectrolyte chains to colloidal core particles leads to spherical polyelectrolyte brushes (SPBs).

Due to the very high concentration of ionic groups in the polyelectrolyte shell, almost all of the corresponding counter ions are confined to the shell. In the presence of monovalent counter ions, this leads to a high concentration of counter ions inside the shell layer. As such, the osmotic pressure of these confined counter ions is very high, leading to a stretching of the polyelectrolyte shell layer. The stretched polyelectrolyte chains evoke the high stability of the SPBs against coagulation, due to both steric (stretching) and electrostatic (charges on the chains) interactions. Addition of traces of multivalent counter ions, however, first results in an ion exchange inside the shell layer, whereby few multivalent ions are getting trapped and many monovalent ions released. The ion exchange process is accompanied by a decreasing particle stability, as the shell layers of the SPBs collapse.



Schematic (left) and cryo-TEM (right) picture of collapsed and unstable SPB particles.

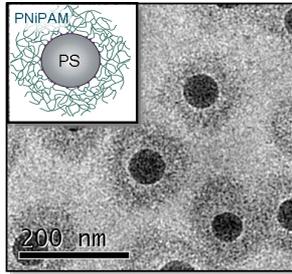
These “electrosteric” stabilized core-shell colloidal systems are widely utilized in industrial applications. We have shown that SPBs serve as good model systems for “electrosteric” stabilized colloids. Thus, we investigate the stability behaviour of SPBs via simultaneous static and dynamic light scattering in the presence of multivalent counter ions.

In the frame of a NSF-DFG cooperation, we work closely with colleagues at the University of California Berkeley and Temple University. During this cooperation, we achieved two goals so far: First, we can now measure the surface potential of an anionic SPB system as a function of the multivalent La^{3+} counter ion concentration in the size order of the thermal energy. Second, given important system parameters of the SPB like contour length and the grafting density of the polyelectrolyte chains, we can predict the stability behaviour of the SPB system within a remarkably high accuracy. Our next steps aim at testing the validity of our model to a wide range of experimental parameters, like SPB-type, type of counter ion and counter ion valency.

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Yielding of a Concentrated Suspension Observed by FT-Rheology: Comparison with the Mode Coupling Theory

Miriam Siebenbürger, Matthias Fuchs*

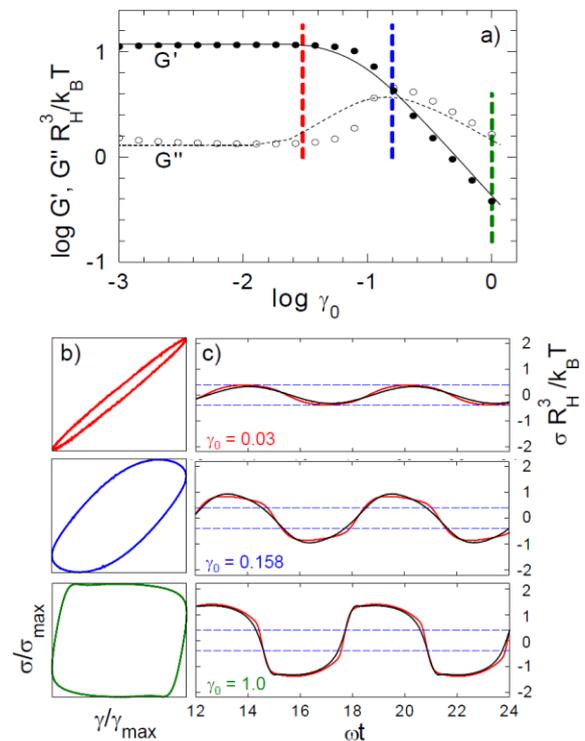


The dynamics and the mechanical properties of hard sphere fluids at high concentrations are an old, but nevertheless a very interesting topic in research. In particular, the transition from the fluid to the glassy state at a volume fraction of $\phi_g \approx 0.58$ and the glass itself above this volume fraction are not completely understood yet. As a model system to mimic the theoretical hard sphere fluid, aqueous suspensions of particles consisting of a poly(styrene) (PS) core and

a thermo-sensitive poly(*N*-isopropylacrylamide) (PNIPAM) shell are used [1]. Suspensions in

the fluid state at high concentrations (close to ϕ_g) show a shear thinning behaviour. In the glass a yield stress σ_y is found. The low force needed for shear melting colloidal glasses and crystals allows rheological investigations at volume fractions above ϕ_g .

The Mode Coupling Theory (MCT) quantitatively describes the dynamics and the mechanical properties of hard sphere suspensions for the fluid and the glassy state close to the glass transition in the stationary flow and the linear viscoelastic regime [2, 3]. The yielding process of the glass can be followed from the linear to the non-linear viscoelastic regime by an oscillatory deformation test as shown in a). The Fourier transformation rheology (FT-rheology) [4] is a quite new rheological technique, which allows the quantification of the degree of non-linearity by means of the intensity ratio of higher harmonics to the fundamental frequency. This quantity can be obtained from the sample by an oscillatory time test at one frequency and strain amplitudes γ_0 as shown in c) by a Fourier transformation. With the schematic MCT-model used to describe the steady state shear and the linear viscoelastic regime it is also possible to calculate the non-linear regime without further parameters (see a) and c)).



- a) Comparison of the experimental deformation test (symbols) at 1Hz in the glassy state at $\phi = 0.65$ with the MCT-calculations (black lines) [5]. At strains of $\gamma_0 = 0.03$ (red), $\gamma_0 = 0.158$ (blue) and $\gamma_0 = 1.0$ (green) time tests with the FT-rheology are performed. The results are given in b) and c).
- b) Lissajous diagrams of the time tests. The enclosed area is directly correlated to the dissipated energy [5].
- c) Time tests at 1Hz and the given strain amplitudes γ_0 ; experimental results are given as black lines. MCT-calculations are drawn in red.

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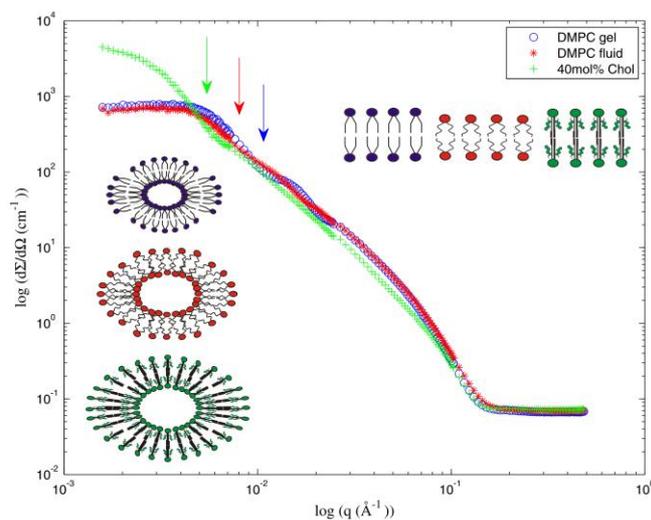
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Fluctuation Dynamics and Elasticity Properties in Unilamellar Phospholipid Vesicles (ULV's): Influence of Temperature and Cholesterol Content

Beate-Annette Brüning

Phospholipid membranes serve as simple model systems to understand basic properties of their far more complex biological counterparts. An important goal in membrane biophysics is to relate the variation of parameters, such as temperature or composition, to changes in structure and dynamics, in order to derive functional properties of an investigated system. In mammal organisms, vesicular membranes often serve as natural carriers (e.g. red blood cells). Mechanical properties of a model membrane and thus the corresponding fluctuation dynamics can be specifically tuned: The insertion of rising amounts of cholesterol causes membrane stiffening, whereas a single lipid membrane softens as its main phase transition between gel and fluid phase is approached upon increasing temperature.

We investigate fluctuation dynamics in unilamellar vesicles and corresponding elasticity parameters under the influence of temperature and cholesterol content using neutron small-angle scattering (SANS) and neutron spin-echo spectroscopy (NSE). The SANS curves in the figure show stable arrangements of single lipid bilayers with diameters between 50 and 60nm (V4, HZB). Characteristic shape fluctuations were investigated in a recent experiment on the long wavelength spin-echo spectrometer IN15 (ILL, Grenoble), and can be described by a normalized intermediate scattering



Small-angle neutron scattering (SANS) curves for unilamellar lipid vesicles (ULV's) obtained on V4, HZB.

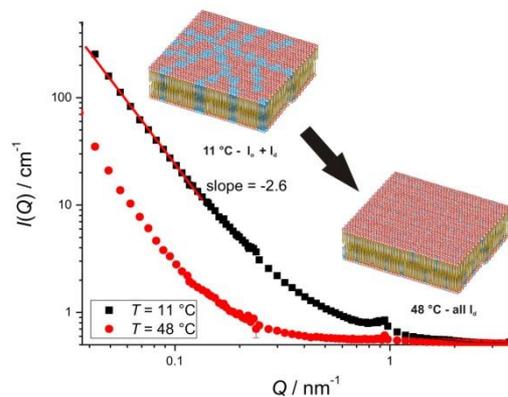
function according to $S(q, t) = \exp(-D_T q^2 t) [A + (1 - A) \exp(-\Gamma_f t)^\alpha]$. For pure bending modes, the relaxation rate follows $\Gamma_f = (k_B T / \kappa)^{1/2} q^3$, thus the bilayer bending rigidity κ is directly obtained, and will be compared with rigidities derived from modeled SANS fits. The recently proposed existence of combined curvature-compression modes will be discussed.

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Small-Angle Scattering on Protein and Lipid Systems

Karsten Vogtt

Proteins and lipids are prominent examples of biological soft matter. Most proteins exhibit their stable, native form in a relatively small temperature window around room temperature. In dilute aqueous solution, they can be treated as colloidal suspensions. In living beings, proteins are involved in a broad range of biological functions, serving as e.g. regulators or as catalysts. Lipids are mainly found as element of structure in the biological cell membranes. Hereby biological cell walls are not “static”, uniform entities separating the cytoplasm completely from its environment, but flexible and dynamic structures which allow local “response” to external stimuli and the controlled exchange of substances over the bilayer. It has been proposed, that local, lateral phase separation into microdomains within the lipid bilayer is involved in the occurrence of such small, functional “patches” - or “lipid rafts”, as they were termed.



Scattered intensity $I(Q)$ of a ternary lipid mixture at two different temperatures. The strong increase of $I(Q)$ at lower temperature is indicative for phase separation with a characteristic spatial mass distribution.

Small angle neutron scattering is an excellent tool to characterize such soft matter systems. Neutrons represent a nearly non-invasive probe, which are sensitive for the atomic nuclei rather than the electron density as it is the case for x-ray scattering. Thus the usually susceptible biological samples are not damaged. Additionally, the sensitivity for different atomic species allows the usage of the so called isotope labelling method. The in biological systems abundant element hydrogen ^1H can be selectively replaced by the isotope deuterium ^2H , which is a much stronger coherent scatterer of neutrons. Thus certain structural patterns in molecular assemblies can be selectively “labelled” and detected. This technique was used to trace phase separation into microdomains in a ternary lipid mixture according to the concept of lipid rafts, as outlined above. The aliphatic side chains of one lipid species were deuterated and small angle neutron scattering was employed to probe for inhomogeneous lateral distribution of lipids within the bilayer as function of temperature (see figure). The results show that at least in such lipid model systems, small microdomains exist on length scales of nanometers and thus would exhibit the appropriate size to act as “functional patches” in biological systems.

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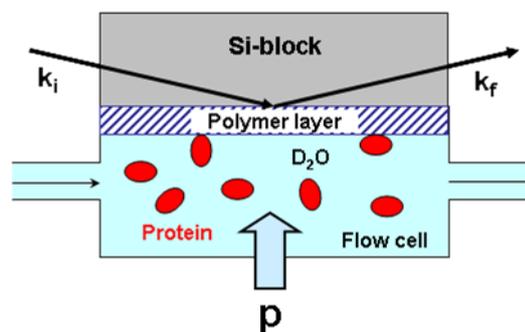
Interfaces: Beamlines and Research

Roland Steitz

The interfaces group takes its name as its mission: The group provides active transport of knowledge between in the interior, the institute of soft matter and functional materials, and the exterior, its short and long term guests and cooperation partners from academia and industry, like a functional unit of a cell membrane.

The interfaces team is responsible for the neutron reflectometer V6 and the new TOF neutron reflectometer BioRef (V18), the latter in cooperation with the Ruprecht-Karls-Universität Heidelberg (Prof. (apl.) R. Dahint, Prof. M. Grunze). The group keeps a strong interactive research profile with its short and long term cooperation partners. Within the Biophysics user laboratory (BioLab) *Interfaces* provides specialized on-site preparation techniques (Layer-by-layer deposition, spin coating, Langmuir-Blodgett and Langmuir-Schäfer deposition) and further off neutron-beamline characterization like ATR-FTIR. In addition we supply our users with complementary x-ray reflectivity and diffraction techniques. Our research activities are committed to studies on structure and functionality of complex interfaces. At present research topics focus on bio-lubrication, responsive solid-liquid interfaces and bio-mimetic systems and their interactions with cellular components as well as on the development of dedicated instrumentation for the purpose.

In 2010 the group served 7 short term external user groups, 5 experimental campaigns from long term cooperation partners, and run 4 in-house sessions on the multipurpose neutron reflectometer V6. Together with our partners R. Dahint and M. Grunze, Ruprecht-Karls-Universität Heidelberg, we are proud to announce finalizing commissioning of BioRef and its first user experiment with C. Garvey from ANSTO, Australia, in August. As a joint activity with RKU the group established routine operation of a dedicated sample cell for neutron reflectivity investigations at solid-liquid interfaces up to 1000 bar hydrostatic pressure. Currently we are working together with partner from the University of Technology Dortmund (TUD), C. Czeslik, in expanding the accessible pressure for neutron reflectivity investigations to even 2500 bar. Together with our cooperation partner on the hard matter side, K. Temst (Catholic University Leuven, Belgium), D. Wallacher from sample environment and colleagues from the institute for complex magnetic materials at HZB, we developed and successfully used a sample cell for simultaneous polarized neutron reflectometry and anisotropic magnetoresistance measurements. A long-lasting and very successful agreement on cooperation on investigations of soft matter interfaces with the Max-Planck-Institute of Colloid- and Interface Science, Golm, ended this year. We are proud to announce that this cooperation agreement was taken up by the Institute of Physical Chemistry, University of Technology Berlin (TUB) with Regine von Klitzing.





Coworkers

Dr. Roland Steitz

Dr. Markus Strobl

Dr. Ralf Köhler

Dipl. Phys. Martin Kreuzer

Dipl. Phys. Matthias Reinhardt

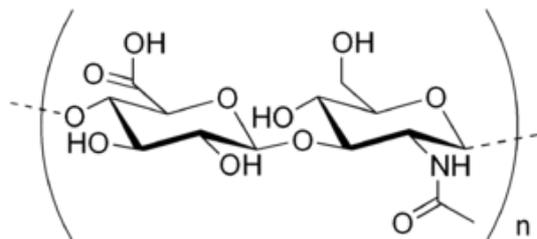
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The Swelling/Stability Effect of Hyaluron on a Lipid Multilayer System

Martin Kreuzer*, M. Reinhardt, M. Strobl, R. Dahint*, R. Steitz



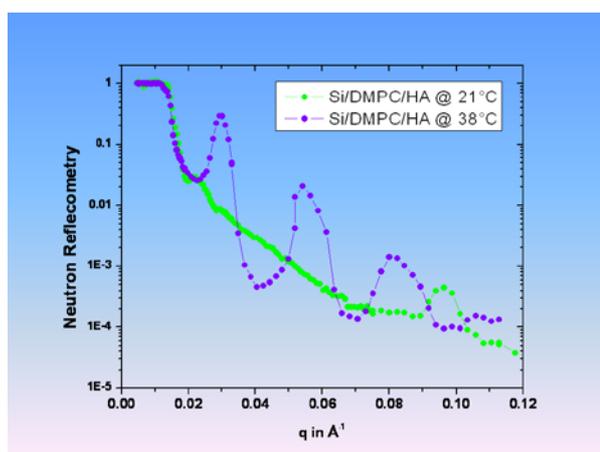
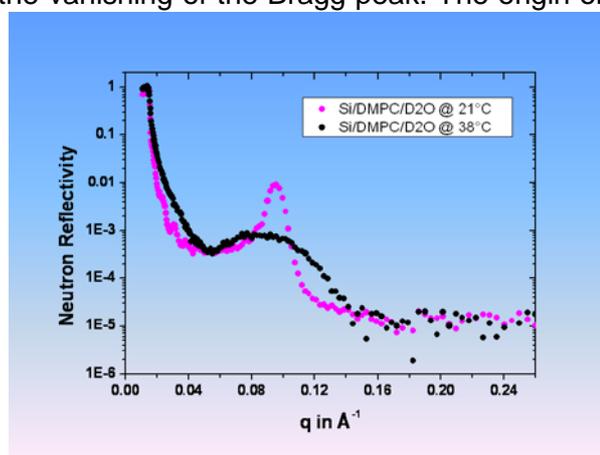
Hyaluron (HA) is a high molecular weight polysaccharide. HA is involved in a wide range of processes in the human body, such as wound healing [1], severe stress, tumor progression and invasion [2]. HA is also known as a lubricant in human joints [3]. Recently we were able to show,

that HA also stabilizes lipid multilayer systems at physiological conditions:

Neutron reflectometry measurements in a flow cell with excess D_2O verified, that a oligolamellar DMPC lipid bilayers coating (bulk phase transition temperature $T_{pt}=23^\circ C$) remained stable on a silicon substrate at $21^\circ C$ in its ordered state (L_β) with a d-spacing of 66\AA , but detached almost completely at $38^\circ C$ in its chain-disordered L_α state from the solid support, in the figure on the right indicated by the vanishing of the Bragg peak. The origin of

the loss of the oligolamellar DMPC bilayer stack at $38^\circ C$ is unclear, but most likely related to the unbinding transition in the chain-disordered state of the lipid lamellae [4]. By contrast oligolamellar lipid bilayers remained stable on a substrate at $38^\circ C$ when incubated with a solution of D_2O with HA: In an independent experiment, carried out at the V6 neutron reflectometer, an oligolamellar lipid bilayers stack was measured against a solution of 3mg/mL HA in D_2O . The sample was investigated shortly after incubating at $21^\circ C$ and after raising sample temperature to $38^\circ C$. The oligolamellar lipid layer remained stable on the substrate, but an immense swelling occurred until a d-spacing of 209\AA , as indicated in the bottom right figure.

In the literature there is no consensus about how a polysaccharide, e.g. HA, affects the lipid phase behavior. The two differing hypotheses are: that (i) the swelling is caused by direct interactions between the solutes and the lipids; or that (ii) the swelling is originated by nonspecific effects related to the osmotic properties of the solutes. Further work is in progress for deriving a clear discrimination and final answer to that open problem.



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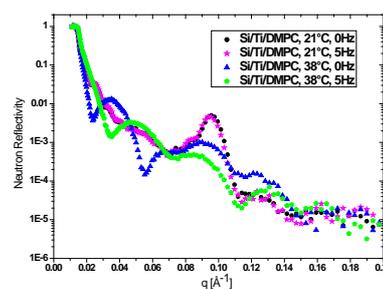
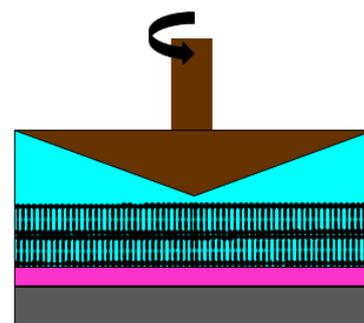
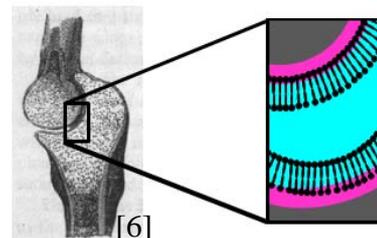
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Lubrication in natural joints – a shear dependence study

M. Kreuzer*, M. Strobl, M. Reinhardt, R. Dahint*, R. Steitz

The search for biocompatible materials has become one of the key issues in modern medicine. Coating of implants by lipid layers is widely used for a better acceptance of the implant in the human organism. With no coverage of their surface artificial implants would invoke macrophage reactions immediately. While on the one hand the durability of implants has been improved significantly, on the other hand the need for permanent and more long-lasting implants is steadily growing [1]. For biomedical applications, titanium-based alloys such as Ti-6Al-4V are most suitable [2]. In case of movable and mechanically stressed implants, such as artificial joints, lubrication under pressure and shear has to be optimized in addition to biocompatibility aspects. While in early artificial joints the movable parts directly contacted each other, researchers nowadays try to copy the principles of lubrication observed in natural joints to reduce friction [3]. Here, the two surfaces of the joint are separated by a liquid phase, the synovial fluid, which mainly contains hyaluronic acid (HA). The most relevant mechanisms and physicochemical parameters to reduce friction are still unclear and subject of controversial discussions. Many studies emphasize the importance of HA for joint lubrication. Furthermore surface-active lipids, which cover the contact areas of natural joints, are considered to play an important role in the reduction of friction [4].



We represented such interface by a suitable model system and employed neutron reflectometry (NR) to study its structural features using a shear setup. The model system was designed as a soft supported lipid membrane (80% POPC + 20% POPS), one bilayer, on the top of a water-swollen polyelectrolyte multilayer on silicon support and incubated in a 3mg/mL solution of HA with D₂O. Our measurements revealed, that an HA-layer of 38Å thickness adsorbs on top of the lipid bilayer. When a shear rate of 2Hz was applied, the HA-layer decreased in thickness to 30Å. Also the lipid membrane thickness decreased from 41Å to 33Å. The same system measured against a pure D₂O did not change its characteristics. Thus, our experiments show that only the combined system of a lipid layer in contact with HA-solution changes, when a shear force is applied.

NR measurements were continued on oligolamellar lipid bilayers directly prepared on a titanium coated silicon substrate, the latter serving as a model for a metallic implant. As seen from the corresponding reflectivity curves, the oligolamellar system became unstable with higher temperature and shear. Further work is needed to clarify the importance of combining of metallic implant surfaces with lipid coatings for forthcoming implant modifications [5].

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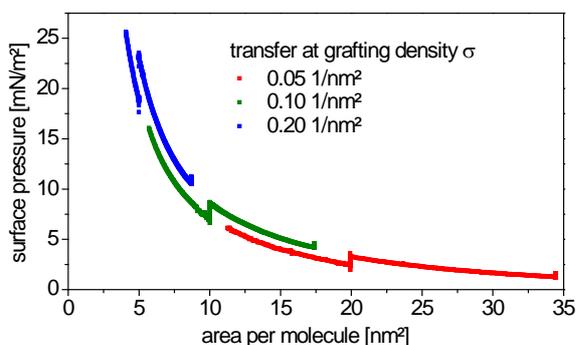
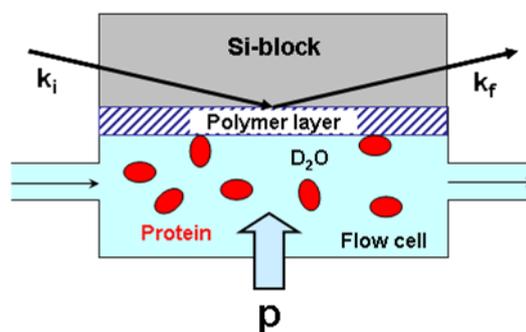
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Functional Interfaces – Brushes and Pressure

Matthias Reinhardt, Martin Kreuzer*, Roland Steitz

High pressure is an important feature of certain natural membrane environments and proteins, for instance in the context of marine biotopes. In that case pressure induced unfolding and denaturation of proteins is of utmost importance. As do natural lipid membranes also polymer brushes provide a soft interface for adsorbed proteins without changing their functionality. We studied adsorbed proteins (BSA) on polymer brushes (dPS-PAA) of different grafting densities at the solid-liquid interface at elevated hydrostatic pressure. Due to the high transparency of solid crystalline materials for neutrons and its high spatial resolution neutron reflectivity (NR) is a perfect tool for investigating structural changes at these solid-liquid interfaces on the nanometer scale when high pressure is applied.

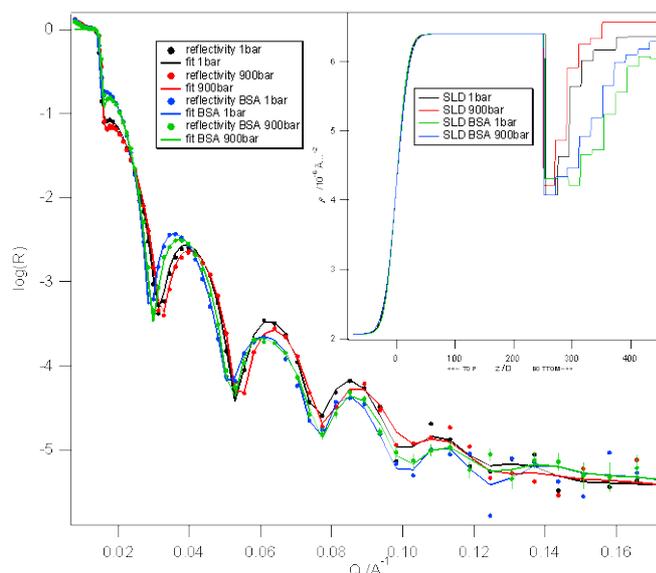


The NR measurements were conducted on V6 at HZB and AMOR at SINQ/PSI.

We found small reversible changes in the reflectivity at elevated pressure. These changes are mainly contributed to an increased SLD of the D₂O subphase. We also found an increased amount of adsorbed BSA proteins for higher grafting densities of the brush.

Elevated hydrostatic pressure up to 900bar does not show significant measurable changes of the adsorbed proteins.

We succeeded in transferring precursor Langmuir films of deuterated poly(styrene)-b-poly(acrylic acid) block copolymers (dPS-PAA) on deuterated poly(styrene) (dPS) pre-coated silicon substrates via Langmuir-Schaefer technique. For this purpose the surface pressure of the precursor Langmuir film provides control of the grafting density and the transfer ratio of the brush as shown in the graph to the left.



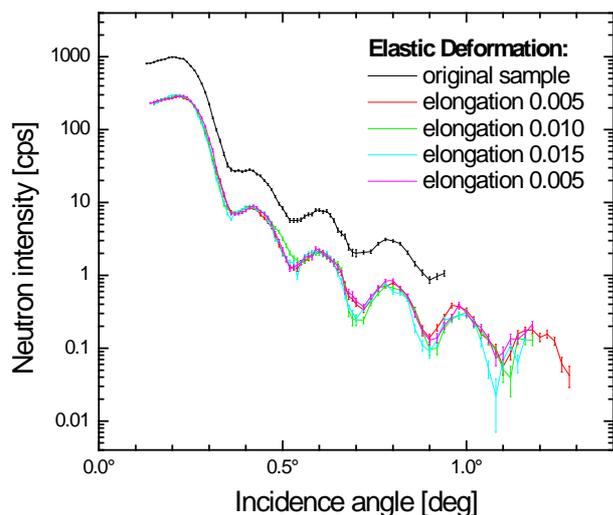
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* Ruprecht-Karls-Universität Heidelberg

Effect of Uniaxial Strain of Polyelectrolyte Multilayers

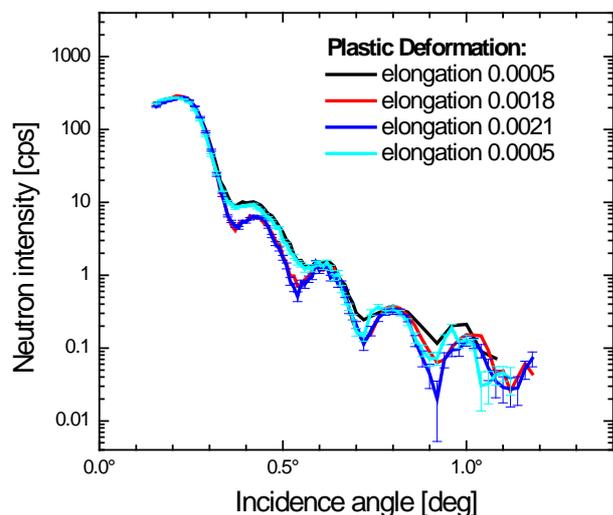
Johannes Früh^{*}, Helmuth Möhwald^{*}, Ralf Köhler[†]

Since their introduction by Decher et al. [1] Polyelectrolyte Multilayers (PEM) have attracted a great scientific interest. The reasons for that are numerous. PEM are easy to prepare on water-born chemistry, the preparation allows for adjusting of thickness and roughness on nanometer scale and with a high reproducibility. PEM are sustainable, a functionalization is possible, and e.g. PEM allow for incorporation of materials (e.g. particles, or even living cells have been incorporated into a PEM matrix) [2].



A key feature for application of PEM is their mechanical behaviour (e.g. stress-strain, or fatigue). Additionally, tests of the mechanics can give information on structure and structural changes of PEM which are not fully understood yet. After having investigated the change of the internal structure for large deformations up to 10% [3], now, we focus on studying effects for small elongations. The aim is to learn about the crossover of reversible and irreversible processes inside the PEM during deformation.

Our new approach is, instead of loading rubber-supported PEM film [3,4,5], to bend a PEM coated thin solid glass substrate. This way a one-dimensional stress is applied [6]. The bent samples are investigated by specular neutron reflectometry whereby the irradiation is perpendicular to the bending axis. Only the specular reflected beam on top of the bent sample is analyzed.



First results were achieved with a PSS/PDADMAC polyelectrolyte system (polystyrene sulfonate / polydiallyldimethyl ammonium chloride) prepared by spraying technique. We found the transition from elastic to plastic behaviour at a very low elongation of ca 0.2%. This is a value typical for solids like aluminium or copper. For polymers one would expect a higher value. Additionally this plastic deformation comes with a slide increase of the film thickness. Both findings are very surprising and seem, at first glance, to

conflict each other. Future studies are planned to clarify these points.

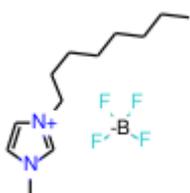
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Anisotropic Fluids at Solid Interfaces: Ionic Liquids

Ralf Köhler^{*}, Rumen Krastev[†], Benilde Saramago[#]

Ionic interactions are the strongest interactions on molecular level, thus materials which are mainly governed by ionic bonds are typically solid at room temperature. Inorganic salts usually also possess ordered crystal lattices. But Ionic Liquids (IL) behave different. They usually have a strong asymmetry in ion size and exhibit, for one ion at least, a relative complex chemical composition. This asymmetry in physico-chemical built-up comes with a broad variety of unusual (macroscopic) properties. The most obvious property was eponymous: Ionic Liquids are liquid salts at room temperature [1,2]. The complex interactions in the ILs yield to complex behaviour, which make IL applicable for many interesting tasks: as lubricants, catalysts, electrolytes (batteries), solvents and dispersants [3,4]. Beside that IL are interesting matters for fundamental science.

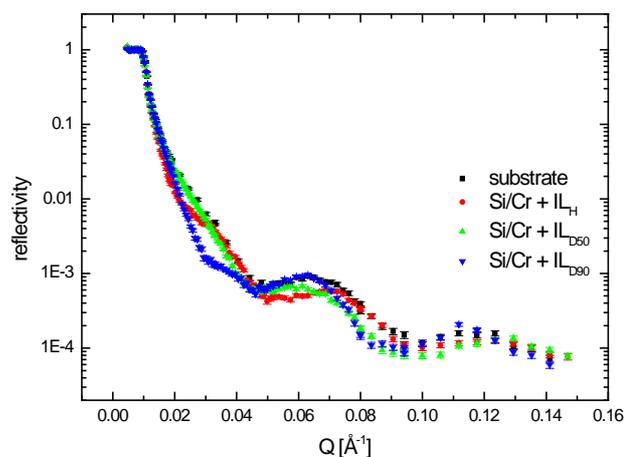


For most of the applications, mentioned above, the interfacial interactions play an important role. This is the motivation to study the wetting behaviour of IL at solid, in this case metallic interfaces, whereby a possible ordering was of special interest. We addressed this topic by using three species of 1-methyl-3-octylimidazolium tetrafluoroborate [OMIM][BF₄] which have a different degree of deuterium (non, 50% and

90%) in the ring of the cationic OMIM-molecule. This labelling would allow for determination of a possible layering of the IL along the OMIM-axis and perpendicular to the surface. Only a few scattering studies address this topic [5,6].

We found evidence for a layered structure in vicinity of the solid surface. The reflectometry curves show different shape although the thickness of the liquid film is almost the same. This study was paralleled by AFM measurements which confirmed the existence of layers parallel to the surface, but, at the same time, gave hints for a more complex structuring at the surface. Most likely ordered and unordered OMIM-layers coexist at different distance towards the solid interface. Further investigations are necessary to clarify

these findings and to allow for establishing a final model of the structural behaviour close to the solid interface.

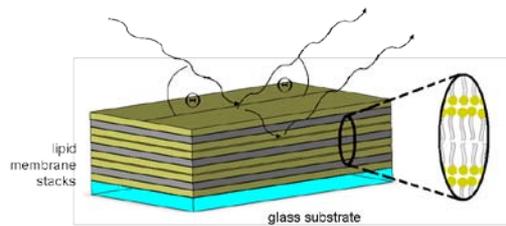


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Influence of Trehalose on the Nano-Structure of Lipid Membranes

M. Gast, U. Fattler*, T. Hauß, H. Haas*, R. Steitz



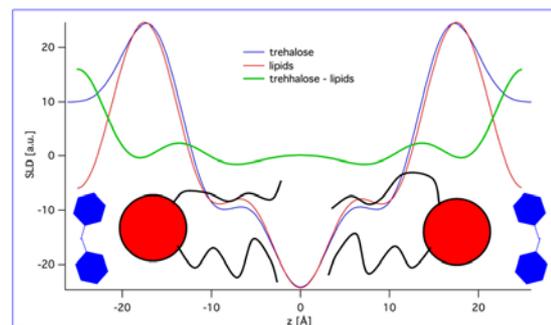
The disaccharide trehalose is known to stabilize cells and their components during drying and freezing processes and it is an important cryo- and lyoprotective excipient in biopharmaceutical manufacturing. For the very reason we investigated the influence of trehalose on the structure of lipid model membranes by x-ray- (XRD) and neutron diffraction

(ND) measurements during which we simulated a drying and rehydration process of the lipid membranes. Our aim was to provide a basis for better understanding and controlling liposome dehydration processes like freeze drying or spray drying.

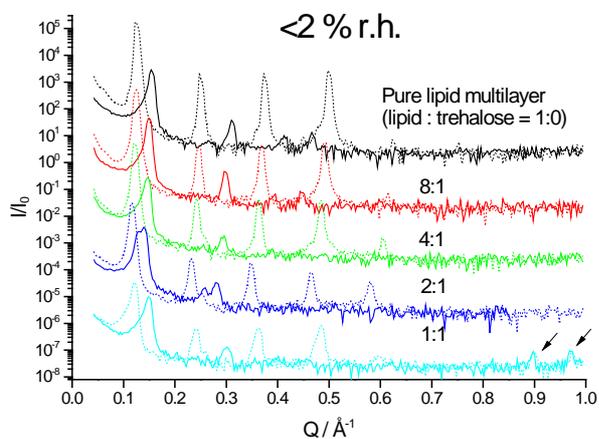
Multilamellar vesicles (MLVs) consisting of 1,2-dioleoyl-sn-glycero-3-trimethyl-ammonium-propane (DOTAP) and 1,2-dioleoyl-phosphatidylcholine (DOPC) were prepared with solutions of different trehalose concentrations, deposited on solid substrates and dried. The obtained stacks were then investigated by XRD and ND under controlled relative humidity (r. h.).

In the presence of trehalose, the lamellar spacing was larger than for pure lipid membranes, indicating that the trehalose inserted in headgroup region of the lipid bilayers.

By neutron scattering measurements with D_2O/H_2O contrast variation we determined the scattering length density profile of the lipid bilayers across their unit cell. From those measurements the localization of water and trehalose molecules was quantitatively deduced: The figure on the right shows that trehalose induced an increase of the SLD in the hydrophilic slab of the bilayers, but it did not penetrate into the lipid bilayer tails region (green line). We found that 2-3 water molecules per lipid headgroup are displaced by trehalose.



X-ray reflectivity measurements (s. below) permitted the illumination of structural implications of dehydrating - rehydrating DOTAP/DOPC membranes in the presence of trehalose. Up to a molar fraction lipid:sugar of 4:1, trehalose was found to insert into DOTAP/DOPC membranes as an integral part of the multilayer stack, which was not excluded from the membrane interface on repeated de- and rehydration. The resulting increase of minimum d-spacing can be considered favourable for preventing liposome fusion or aggregation.



Full reversibility of structural reorganizations as a function of relative humidity thus nominates DOTAP/DOPC liposomes in trehalose robust regarding dehydration protocols in the course of manufacturing and application of pharmaceutical DOTAP/DOPC liposome products.

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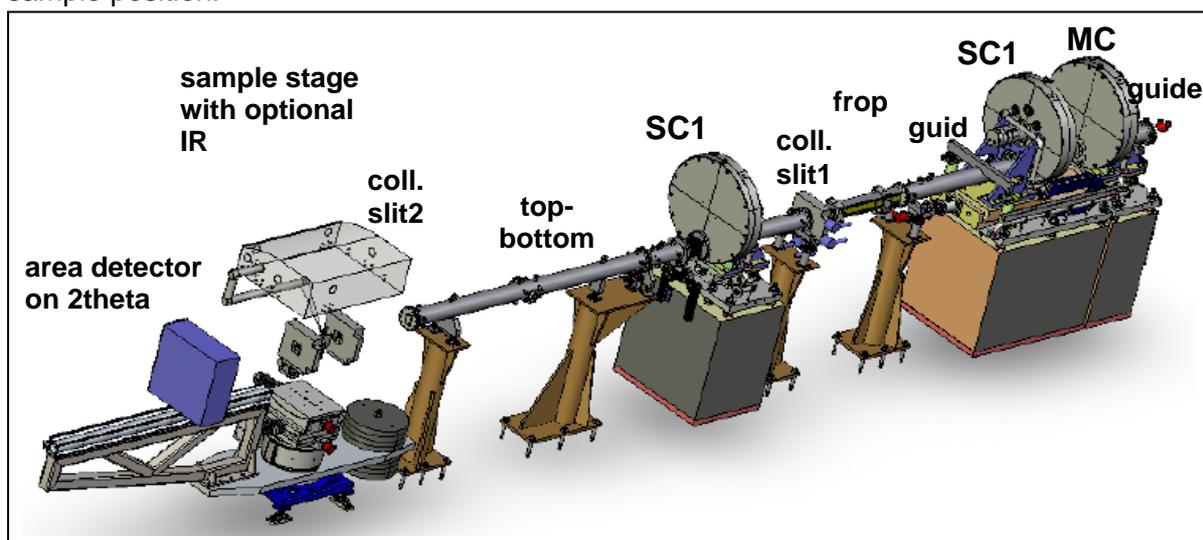
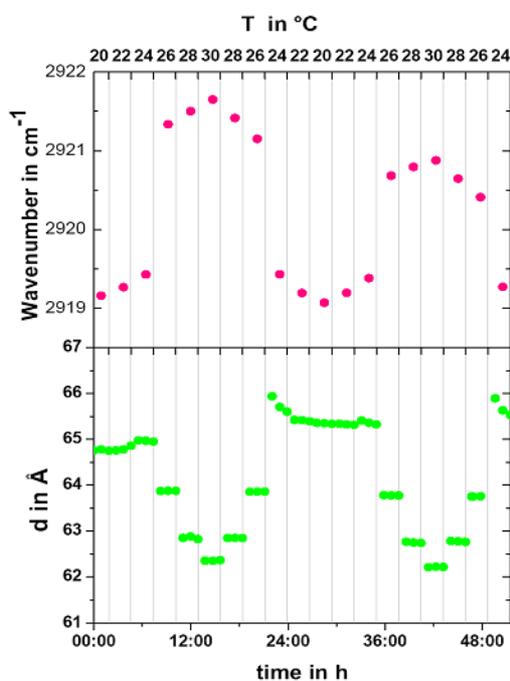
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* Medigene AG, Martinsried

V18 BioRef – a Versatile Tool for Surface/Interface Characterizations

Markus Strobl, Martin Keuzer*, Reiner Dahint*, Roland Steitz

BioRef is a time-of-flight neutron reflectometer, has recently been realized in the framework of a BMBF funded project in cooperation with the University of Heidelberg. The instrument, which became operational early 2010 is mainly dedicated to investigations of solid-liquid soft matter surfaces and interfaces, which serve as model systems of Biological systems. For this purpose sample environment capable to mimic physiological conditions in terms of temperature, flow and shear as well as pressure has been and is further developed. Besides the flexibility of the instrument, which allows for tailoring instrumental conditions, like resolution and utilized wavelength band to the specific requirements of investigations and hence to enable even kinetic studies in selected scattering vector ranges, the set-up offers the unique option of in-situ attenuated total reflection Fourier transformed infrared (ATR-FTIR) spectroscopy. Such spectroscopy can be utilized to gain conformational information complementary to the structural data concerning the scattering length density (SLD) profiles deduced from the neutron reflectivity. This way e.g. the unfolding of proteins which indicates a loss of their functionality at investigated surfaces can be observed under the very same conditions under which the structural data is collected even when such conditions are not constant with time, i.e. under kinetic conditions. A first combined neutron-IR study of a lipid multilayer during a temperature scan is presented in the Figure on top. The Figure on the bottom of the page is a drawing of the principal instrument layout featuring the IR spectrometer on top of the sample position.



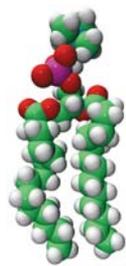
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[2] M. Strobl, R. Steitz, M. Kreuzer, M. Rose, H. Herrlich, F. Mezei, M. Grunze and R. Dahint, *Review of Scientific Instruments* **2011**, submitted.

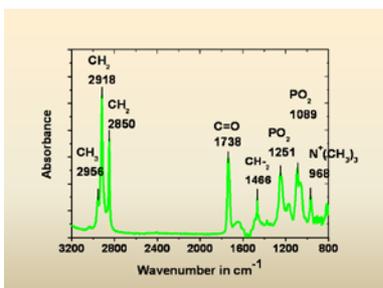
* Ruprecht-Karls-Universität Heidelberg

Fourier Transform Infrared spectroscopy (FTIR) at Solid-Liquid Interfaces

Martin Kreuzer*, Marie Charlotte Hemmer, Reiner Dahint*, Roland Steitz



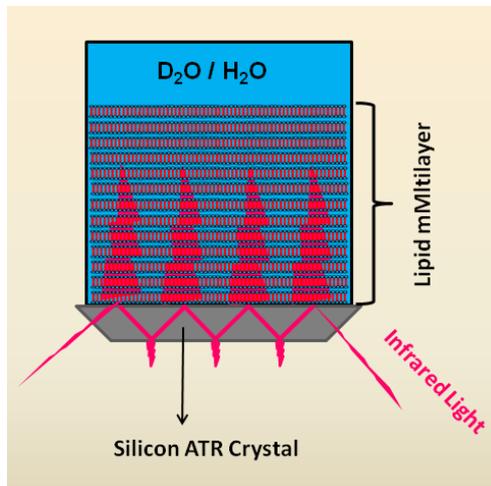
Due to their hydrophilic head and hydrophobic tail groups lipid molecules are able to form multilayer membranes in an aqueous environment. Up to 40% of the molecules in cell membranes are lipids [1]. These biological membranes are essential for directed and proper molecular life functions [2]. Fundamental for understanding the functional properties of membrane lipids is a detailed knowledge of their preferred molecular conformations. Attenuated total reflectance (ATR) - FTIR is the favored technique for examining directly multilayer membranes in an aqueous environment. We utilized a BioATR II setup from Bruker Optics, where an infrared beam gets reflected several



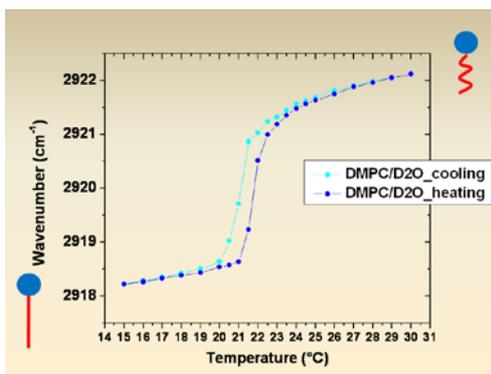
times at a silicon-water interface,

before it is detected with a nitrogen cooled, mercury cadmium telluride (MCT) detector. The infrared absorbance signal from a lipid multilayer, attached to the silicon surface and incubated with an aqueous solution, is enhanced with

every reflection at the interface. In addition, the temperature controlled setup made it possible to measure the confirmation of lipid membranes in a wide range of temperatures against different aqueous solutions. In particular the CH₂ vibrations of the tail groups of the lipids gave information about their lamellar phase (e.g. gel-like P_β or fluid-like L_α phase) [3].



enhanced with



The temperature dependent measurements in the ATR-FTIR setup with a lipid multilayer against excess D₂O showed a shift of the symmetric and anti-symmetric CH₂ stretching band at the phase transition. For the lipid molecule 1,2-Dimyristoyl-sn-Glycero-3-phosphocholine (DMPC) the symmetric CH₂ stretching band shifted from 2850cm⁻¹ to 2853cm⁻¹, corresponding to the P_β and L_α phase, respectively, when the temperature was scanned between 15°C and 30°C. The asymmetric CH₂ stretching band shifted from 2918cm⁻¹ to 2923cm⁻¹. The most pronounced shift occurred at 21.5°C, which is close to the phase transition of the bulk lamellar phase of DMPC in H₂O at app. 24°C.

The design of an advanced ATR-FTIR setup including a sample cell made it possible to perform a combined measurement of Neutron Reflectivity and ATR-FTIR on the same sample, at the same time. This unique setup is realized at the BioRef neutron beamline at the HZB. Here, conformational as well as structural information of the interface can be measured.

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* Ruprecht-Karls-Universität Heidelberg

Pressure cell for investigations of solid-liquid interfaces by neutron reflectivity

Martin Kreuzer*, Thomas Kaltofen*, Beat H. Zehnder#, Reiner Dahint*, Roland Steitz

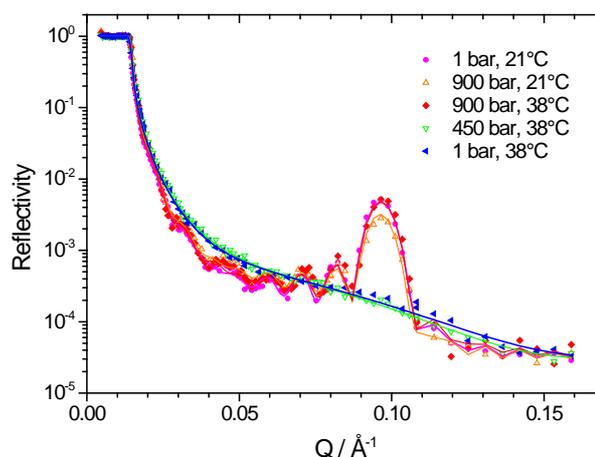
Studies of the high pressure phase behaviour of lipid and surfactant systems, in particular of phospholipid bilayers, which can serve as model biomembranes, have become a standard setup in bulk systems.^[1] Sample cells for hydrostatic pressure conditions up to 7000 bar for use with neutron and X-ray diffraction techniques are available nowadays.^{[2], [3]} Design and manufacturing of those cells is favoured by the small sample volume, typically 0.04-0.5 mL, needed for the diffraction experiments. Cross sections of these pressure cells are of the order of a few mm² and thus easy to handle up to highest pressures. Reflectivity studies on confined systems and even single lipid membranes immobilised at



solid-liquid interfaces on the opposite require large surface areas typically of the order of some tens of cm² and some tens of mL liquid volume. These pre-conditions hindered complementary investigations of surface bound systems for very long.

We recently succeeded in developing an apparatus for measuring scattering length density and structure of molecular layers at planar solid-liquid interfaces under high hydrostatic pressure conditions (figures).^[4] The device is designed for in situ characterizations utilizing

neutron reflectometry in the pressure range 1–1000 bar at temperatures between 5 and 60 °C. The pressure cell is constructed such that stratified molecular layers on crystalline substrates of silicon, quartz, or sapphire with a surface area of 2800 mm² can be investigated against noncorrosive liquid phases. The large substrate surface area enables reflectivity to be measured down to 10⁻⁵ (without background correction) and thus facilitates determination of the scattering length density profile across the



interface as a function of applied load. Our current interest is on the stability of oligolamellar lipid coatings on silicon surfaces against aqueous phases as a function of applied hydrostatic pressure and temperature but the device can also be employed to probe the structure of any other solid-liquid interface.

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* Ruprecht-Karls-Universität Heidelberg

SITEC-Sieber Engineering AG, Ebmatingen

Colloid Chemistry

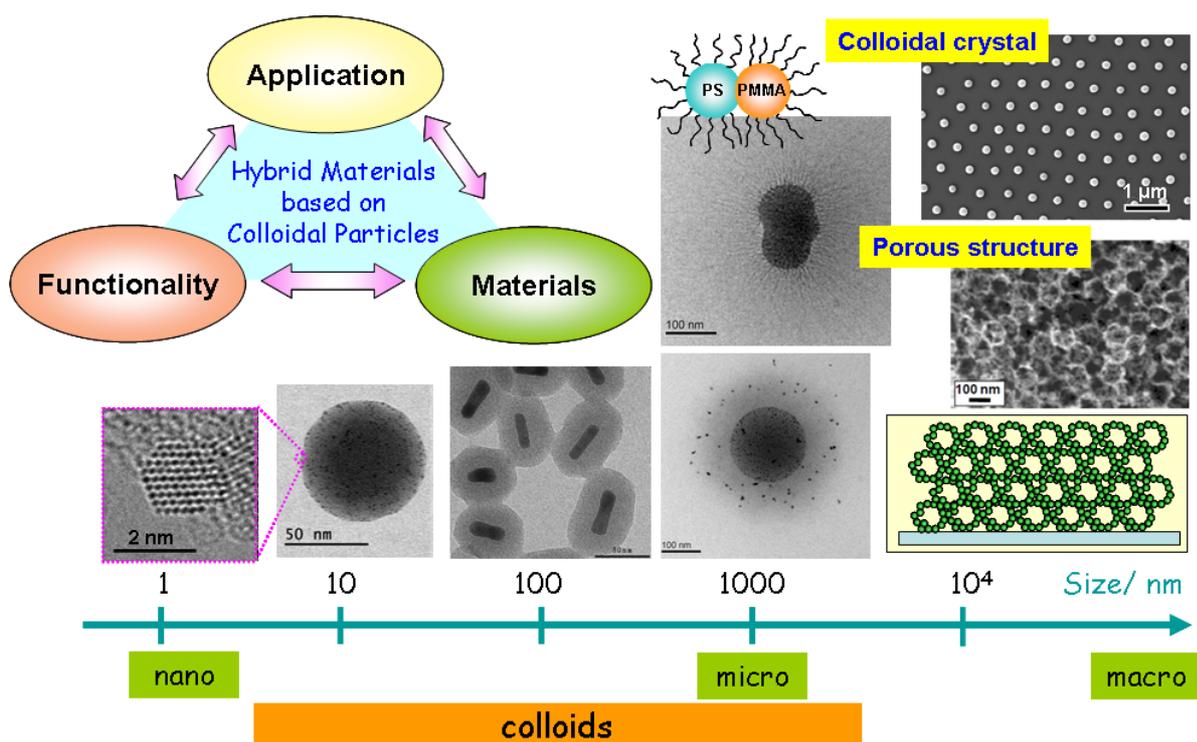
Yan Lu

The research work in the Colloid Chemistry Group mainly focuses on the design and fabrication of functional hybrid materials based on colloidal particles with versatile applications, such as catalysts, solar cells, and optical devices.

For this purpose, composite particles based on metal nanoparticles (Au, Ag, Pd, Pt, etc.), metal nanoalloys (Au-Pt, Au-Pd, etc.) and metal oxide (TiO_2 , ZrO_2 , MnO_x , etc.) particles using colloidal particles as carrier system have been prepared. Compared to other reported carrier systems, colloidal particles have various merits, such as superior stability, facile synthesis for industrial potential, good control over particle size and composition, and easy functionalization providing novel properties. Thus, these composite particles have multiple functionalities with improved physical and chemical properties in a feasible way. These nanocomposite particles have been proven as excellent (photo)catalysts for various chemical catalytic reactions that proceed in aqueous solutions or in two-phase systems. Kinetic studies of catalytic reactions are an essential part of our research in order to understand the mechanism of the reaction in the presence of metal nanoparticles.

Special efforts have been made for the preparation of anisotropic particles, which show interesting properties in light scattering and plasmon absorption. For example, dumbbell-shaped colloids with a size around 200 nm can be prepared by seeded-emulsion polymerization, which can be used as core for the further deposition of well defined water soluble polyelectrolyte brushes or stimuli-responsive shell. On the other hand, hybrid structures based on Au nanorods can be applied as model system to study the plasmon effect of metal nanoparticles on the kinetics and efficiency of photoelectrocatalysis. In addition, possible applications like a surface plasmon polariton laser will be investigated.

We expect that the full knowledge of controlled synthesis of colloidal particles will benefit our users. Moreover, cooperation with research groups in the field of solar cells, plasmonics and catalysis within and outside HZB will help us not only to expand the application spectrum of composite particles but also result in vital understanding of the fundamentals.



Coworkers:

Dr. Yan Lu

Dr. Martin Hoffmann

Dipl Chem Frank Polzer

Dipl.Chem. Julian Kaiser

Dipl.Chem. Stefanie Wunder

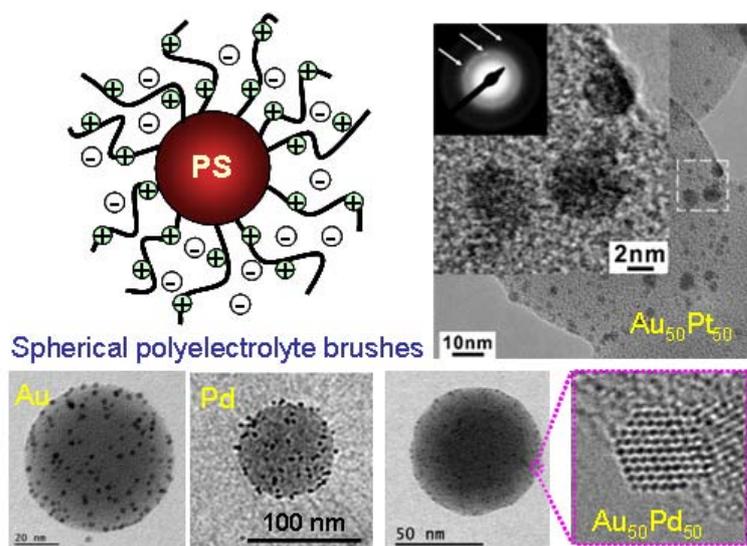
M.Sc. Shuang Wu

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5. S. Wunder, F. Polzer, Y. Lu, M. Yu, M. Ballauff, "Kinetic Analysis of Catalytic Reduction of 4-Nitrophenol by Metallic Nanoparticles immobilized in Spherical Polyelectrolyte Brushes", *J. Phys. Chem. C* **2010**, 114 (19), 8814-8820.
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8. J. Yuan, F. Schacher, M. Drechsler, A. Hanisch, Y. Lu, M. Ballauff, A. Mueller, "Stimuli-Responsive Organo-Silica Hybrid Nanowires Decorated with Metal Nanoparticles", *Chem. Mater.* **2010**, 22, 2626-2634.
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Spherical Polyelectrolyte Brushes as “Nanoreactors” for Metal Nanoparticles or Nanoalloys

Julian Kaiser, Yan Lu, M. Albrecht (Leibniz-Institut für Kristallzüchtung, Berlin)



Metal nanoparticles and nanoalloys have been raised a lot of interests in the present. The biggest problem is the prevention of metal nanoparticles from agglomeration. During our study, spherical polyelectrolyte brushes (SPB) have been used as “nanoreactors”, in which metal nanoparticles can be immobilized and handled in an easier fashion. The SPB particles consist of a polystyrene core, onto which long

linear chains of polyelectrolyte are densely grafted. Metal ions will be immobilized into the surface layer of polyelectrolytes of the SPB as counterions. Reduction of these immobilized metal ions with NaBH_4 leads to nanoparticles of the respective metal. During our study, various nanoparticles of noble metals (such as Au, Pd, Pt, etc.) can be immobilized in this way and used for catalysis in aqueous media, that is, under very mild conditions. Thus, the composite systems of metallic nanoparticles and spherical polyelectrolyte brushes allow us to do “green chemistry” and conduct chemical reaction in a very efficient way.

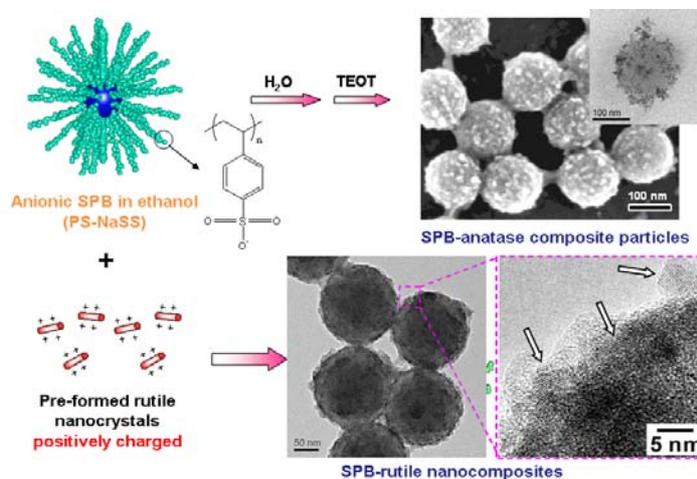
Moreover, metal nanoalloys have attracted more interests than metal nanoparticles recently due to the fact that their chemical and physical properties may be tuned by varying the composition and atomic ordering, which are important in determining chemical reactivity and especially catalytic activity. During our research, we have demonstrated that spherical polyelectrolyte brushes (SPB) can work efficiently as a carrier system for the immobilization of metal nanoalloys (such as Au-Pt, Au-Pd). In this case, with the help of high resolution transmission electron microscopy (HR-TEM), powder x-ray diffraction (XRD) and extended x-ray absorption fine structure (EXAFS), it is possible for us to get precise information about the dynamics and compositional and structural evolution of these metal nanoalloy particles. In addition, the catalytic activity of the generated nanoalloys has been tested by the reduction of 4-nitrophenol. The rate constant normalized to metal surface area goes through a maximum for the catalytic reduction as the function of Au amount in bimetallic nanoparticles, which indicates a synergistic effect of both metals related to their catalytic activity.

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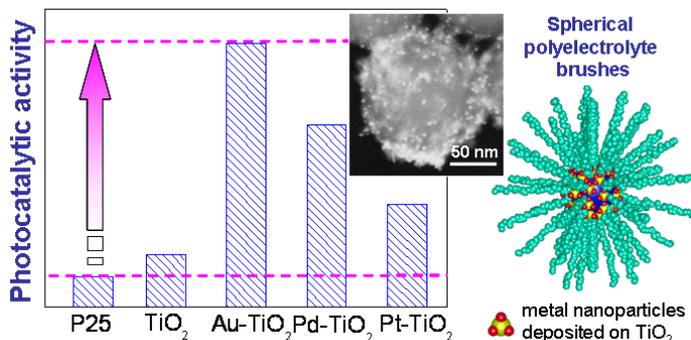
Spherical Polyelectrolyte Brushes as “nanoreactor” for Well-Defined Crystalline TiO₂ Nanoparticles

Yan Lu

TiO₂ nanomaterials have received much attention recently due to their photocatalytic activity, high chemical stability and possible applications in solar cells. Moreover, mesoporous TiO₂ networks with high surface area are of particular interest for a number of applications. Colloidal latex particles have been used for the preparation of hollow TiO₂ spheres or continuous macroporous TiO₂ structures. However, all as-prepared TiO₂ composites prepared in this way by a sol-gel approach are amorphous. The latex particles act only as a template for the macroporous structure and calcination is required to achieve sufficient crystallinity.



Spherical polyelectrolyte brushes (SPB) particles may serve as well-defined nanoreactors for the immobilization of TiO₂ nanoparticles. The SPB particles consist of a solid PS core from which long anionic polyelectrolyte chains are densely grafted. Crystalline anatase or rutile TiO₂ nanoparticles can be generated directly by “sol-gel” method or electrostatic adsorption



in the presence of SPBs at low temperature, respectively. Thus, composite particles of a polymeric carrier and crystalline TiO₂ in a well-defined modification can be obtained without any further heat treatment. In addition, the as-prepared TiO₂ nanocomposites exhibit an excellent colloidal stability. The photocatalytic

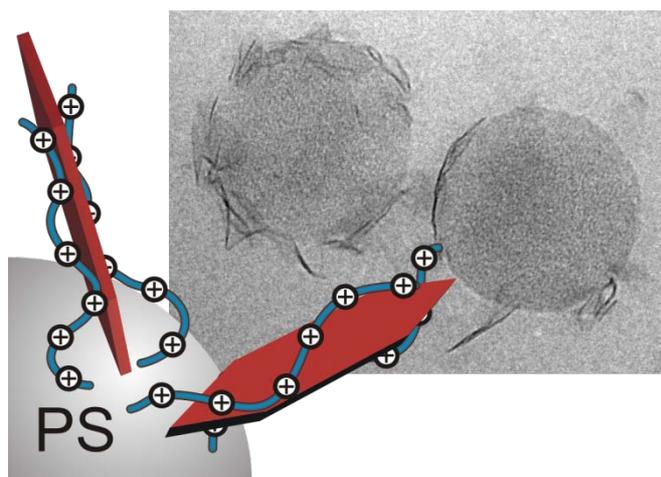
activity of the composite particles for the degradation of dye RhB under UV irradiation can be dramatically enhanced after deposition of metal nanoparticles on it. Finally, calcination of the composite particles leads to a macroporous scaffold of mesoporous TiO₂ nanoparticles, which are thermally stable against collapse. Possible applications, as e.g. for solar cells, have been proved.

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Formation of Ultrathin Birnessite-Type Nanoparticles Immobilized on Cationic Spherical Polyelectrolyte Brushes

Frank Polzer

Layered manganese oxide materials attracted much interest because of their potential applications as catalysts and electrode materials. In both fields, high surface to volume ratios are favored. For layered materials, multistep processes like intercalation of bulky ions with subsequent delamination are needed to obtain thin- or monolayered structures with high surface areas.



We present a new method of in situ formation and stabilization of ultrathin, layered manganese oxide nanoparticles (MnO_xNP) in aqueous solution by using spherical polyelectrolyte brush particles (SPB) without any further reducing agent and delamination procedure.

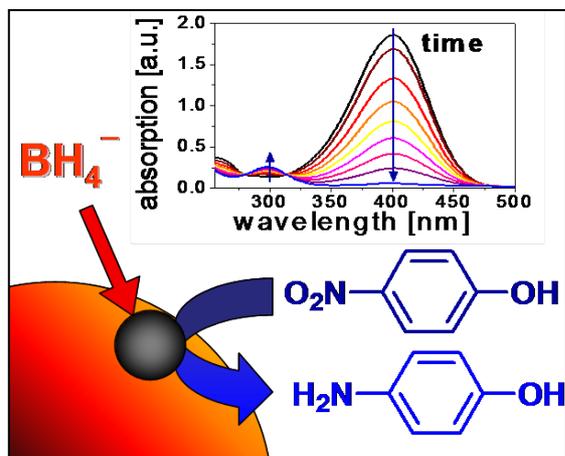
A combination of TEM, cryoTEM, PXRD, EDX and XAS studies was used for detailed characterization. The

nanoparticles are of birnessite type, a layered material composed of hydrous lamellae of hexagonal MnO_x sheets. These sheets form nanoparticles with an average length of 20 nm and a breadth of ca. 1.6 nm indicating a composition of single lamellae or of ultrathin stacks of very few lamellae. The individual layers have a stacking disorder leading to hk bands in the PXRD pattern which is typical for c^* -disordered K^+ birnessite. XAS proved to be an excellent way to gain important information on the poorly crystalline nanoparticles such as the interatomic distances, the coordination chemistry and the average oxidation state. Furthermore, the nanoparticles are well stabilized against coagulation by immobilization onto the SPB carrier particles in aqueous solution. This finding can be traced back to a strong interaction the positively charged poly(trimethyl ammonium ethyl methacrylate chloride) pTMAEMC chains of the SPB with the negatively charged birnessite particles. First catalytic studies show great potential for a use as an oxidation catalyst for alcohols and for epoxidation reactions.

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Kinetic Studies of Reduction of 4-Nitrophenol Using Metal Nanoparticles immobilised in Spherical Polyelectrolyte Brushes as Catalyst

Stefanie Wunder, Yan Lu



Metallic nanoparticles (NP) have been the subject of intense research during the recent years because of their potential use in catalysis. A model reaction suitable for this purpose should be well-defined, that is, no by-products should be formed. Moreover, the degree of conversion should be easily monitored by a simple and fast technique. Therefore the catalytic reduction of 4-nitrophenol (Nip) to 4-aminophenol using sodium borohydride (BH_4^-) as reducing agent has been chosen. The kinetic of the reaction

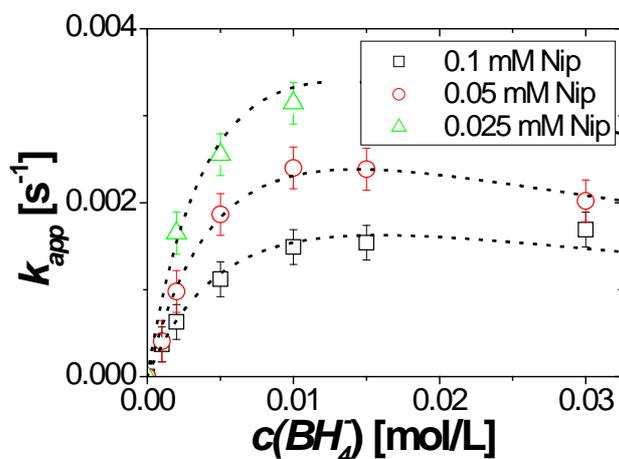
can be analyzed by UV-vis spectroscopy. From the time-dependency of the adsorption peak of Nip at 400 nm the apparent rate constant k_{app} can be calculated.

The proposed reaction mechanism comprise, that both educts must adsorb onto the surface of the metallic nanoparticles. This mechanism is also called the Langmuir-Hinshelwood model:

$$k_{app} = \frac{k \cdot S \cdot K_{Nip}^n \cdot c_{Nip}^{n-1} \cdot (K_{BH_4} \cdot c_{BH_4})^m}{(1 + (K_{Nip} \cdot c_{Nip})^n + (K_{BH_4} \cdot c_{BH_4})^m)^2}$$

Furthermore by varying the concentrations of the educts (Nip and BH_4^-) and applying the Langmuir-Hinshelwood reaction model, the reaction rate k and the adsorption constants of 4-nitrophenol K_{Nip} and sodium borohydride K_{BH_4} , can be determined for different metallic nanoparticles, respectively. Another

measured variable of this model reaction is the induction time. We could show that it is highly dependent on the concentration of Nip but not on that of BH_4^- . Hence, we assume that first a surface restructuring process must occur, before the reaction can start. Further investigation will be focuses on the influence of temperatures on the reaction kinetics in order to get activation energy and adsorption enthalpy by using the van't Hoff equation of the reaction. In addition, it will be also interesting to combine this part of work with theoretic simulation.



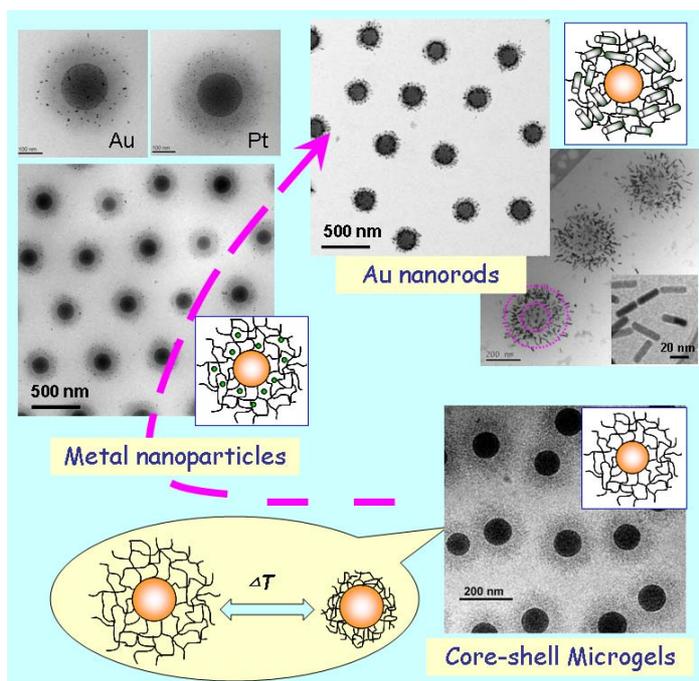
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Thermosensitive Core-Shell Microgels as Active “Nanoreactors” for Metal Nanoparticles

Yan Lu



Metal nanoparticles have attracted much attention because such particles may exhibit properties differing strongly from those of the bulk metal. However, the metallic nanoparticles must be stabilized in solution in order to prevent aggregation. In principle, suitable colloidal carrier system may be used as a “nanoreactor”, in which the metal nanoparticles can be immobilized and used for the purpose at hand. The use of microgel particles as reactors for the deposition of metal nanoparticles may have several important advantages over other systems,

namely, stability, ease of synthesis and easy functionalization providing stimulus-responsive behavior.

Thermosensitive core-shell microgel particles, in which the core consists of poly (styrene) (PS) whereas the shell consists of poly (N-isopropylacrylamide) (PNIPA) network crosslinked by N, N'-methylenebisacrylamide (BIS), can be used as “nanoreactors” for the deposition of catalytically active metal nanoparticles. Different metal nanoparticles (such as Ag, Au, Pd, Pt and Rh) as well as Au nanorods can be homogeneously embedded into the microgel particles.

The catalytic activity as well as the optical property of metal nanoparticles immobilized in thermosensitive microgels can be tuned by the volume transition within the microgel. At low temperatures, the composite particles are suspended in water, which swells the thermosensitive network attached to the surface of the core particles. At higher temperatures ($T > 32^{\circ}\text{C}$), the PNIPA-network, however, undergoes a volume transition, in which most of the water is expelled. We demonstrate that the catalytic activity of the microgel-metal nanocomposites can be tuned by the volume transition within the microgel of these systems by using the catalytic reduction of 4-nitrophenol as the model reaction. Thus, the microgel particles present an “active” carrier system for applications in catalysis.

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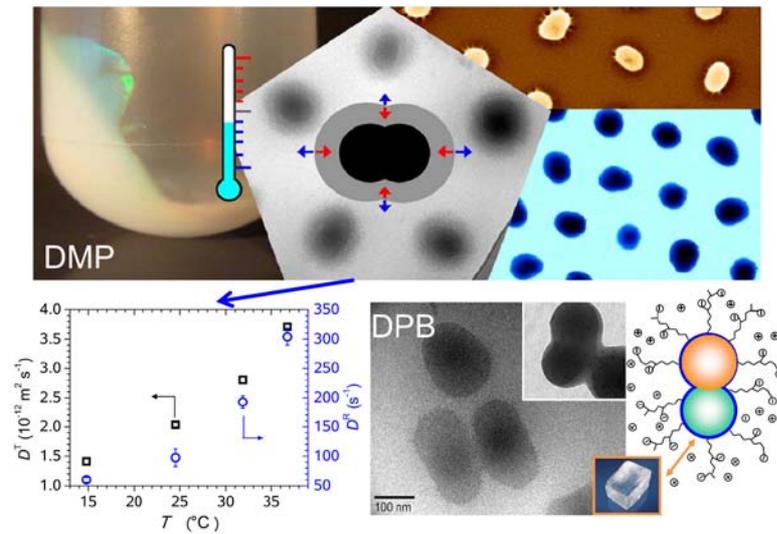
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Dumbbell-Shaped Core-Shell Nanoparticles

Martin Hoffmann



Spherical colloidal particles have been thoroughly investigated for the last decades and they are still used as model systems in many theoretical studies. However, the development of anisotropic colloidal particles with simple or complex shape and their dynamics in solution are fundamental towards an understanding of many problems like sedimentation,

coagulation or rheology. Several strategies are available to obtain rigid dumbbell-shaped colloids with a size ranging between 100 nm and several microns. Potential applications of these structures demand high stability and an easy modification of the particle morphology. The combination of a rigid dumbbell-shaped polymer core with a well defined water soluble stimuli-responsive shell may fulfill these requirements.

With this aim, two systems (DPB / DMP) were prepared and their morphology was investigated by means of electron microscopy techniques. Depolarized dynamic light scattering was used to follow the dynamics in solution since the particles depolarize light. For system I, a dense layer of polyelectrolyte chains was attached to the polymer core particle which allows to tune the particle size and the aspect ratio by the ionic strength as external stimulus ("DPB"). System II consists of dumbbell-shaped polymer cores which carry a thermoresponsive shell made of a poly(*N*-isopropylacrylamide) network crosslinked by *N*, *N*-methylenebisacrylamide ("DMP").

For the DPB, the condensation of the counterions to the polyelectrolyte chains leads to a high osmotic pressure within the brush and thus to an almost fully extended shell in water. Adding salt screens the electrostatic interactions in the brush and leads to the shrinking of the shell. In the case of the DMP, the thickness of the shell is determined by the solubility of the thermoresponsive network in water. We demonstrate that the translational (D^T) and the rotational diffusion (D^R) of the particles as measured by depolarized dynamic light scattering significantly depend on the shell thickness. For the future, the DMP present a versatile model system to investigate the fluid-solid transitions of concentrated dispersions of both thermoresponsive and anisotropic colloids.

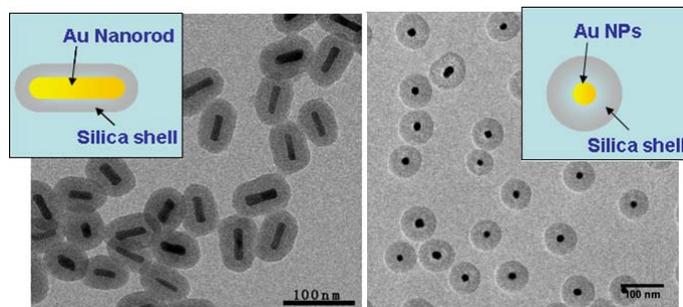
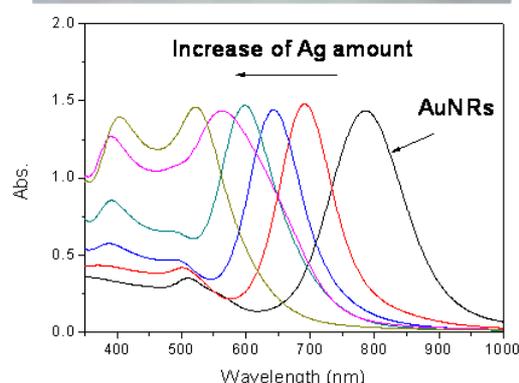
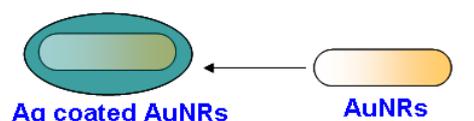
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Plasmonic Au – Based Nanocomposite Particles: Synthesis, Characterization and Application

Shuang Wu, Yan Lu, Oliver Benson*, Thomas Aichele*

Plasmonics has become one of the most active fields in nanophotonics. In last several years, there has been a rapidly increasing activity within this field as its wide application field ranging from sensing and biomedicine to imaging and information technology. Previous reports have demonstrated that the shape and structure of metal nanocrystals play the most important roles in determining the number, position, and intensity of localized surface plasmon resonance modes.

During our research work, Au and Ag nanocrystals with defined size and shape have been fabricated by the simple wet-chemical method, which is one of favoured routes toward the cost-effective large scale production of metal nanostructures. For example, gold nanorods (AuNRs) can be prepared by seeded-mediated method, which provide two intrinsic (transverse and longitudinal) bands. Deposition of Ag on the AuNRs surface leads to a blue shift in the longitudinal surface plasmon absorption band of Au. Thus, it is possible to prepare nanoparticles with controllable surface plasmon band by this approach.



On the other hand, surface plasmon polaritons (SPP) can guide light and confine it to sub-wavelength dimensions. Recently, evidence for a surface plasmon laser (spaser) was reported. The relatively large loss in the metallic spaser cavity can be compensated by significant gain in a surrounding gain material. Thus, it is

crucial to design a metal core/conjugated organic shell hybrid structure that shows coherent amplification of SPP modes and spaser action. For this purpose, metal nanoparticles with different shapes will be synthesized. After covering these metal nanoparticles with a homogeneous SiO_2 layer, organic molecules will be immobilized inside to form the metal core/conjugated organic shell hybrid structure. Here, developing novel metallo-organic nanoparticles is essential not only for the practical applications but also for fundamental understandings.

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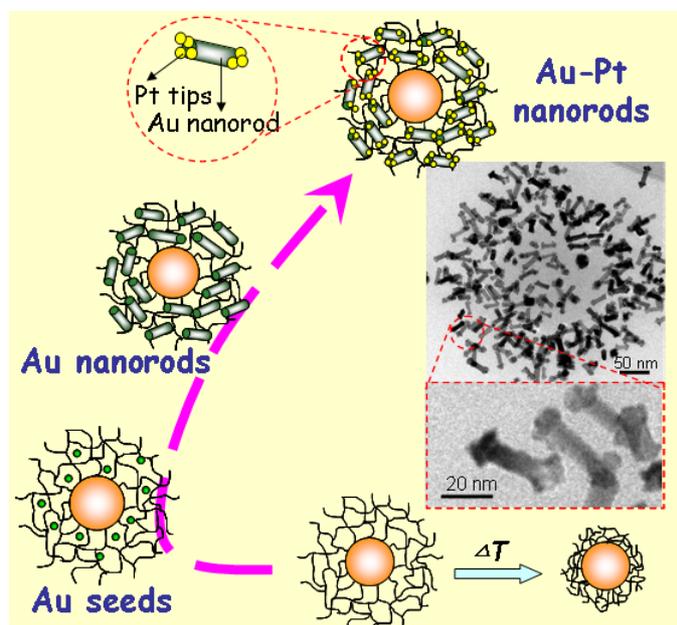
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In-situ Growth of Catalytic Active Au-Pt Bimetallic Nanorods in Thermo-Responsive Core-Shell Microgels

Yan Lu, J. Yuan*, F. Polzer, M. Drechsler*, J. Preussner†

Bimetallic Au-Pt nanorods (NRs) can be *in-situ* grown into thermosensitive core-shell microgel particles by a novel two-step approach. This demonstrates for the first time that the control of the shape as well as hierarchical structure of nanoparticles in the microgel particles is achievable via an *in situ* strategy. The volume transition of microgel network leads to a strong red shift of the longitudinal plasmon band of the Au NRs, which are immobilized in microgel networks.



Platinum can preferentially deposit onto the tips of Au NRs to form dumbbell-shaped bimetallic nanoparticles. The novel synthesis forms bimetallic Au-Pt NRs immobilized in microgels without impeding their colloidal stability. Quantitative analysis of the catalytic activity for the catalytic reduction of 4-nitrophenol indicates that bimetallic Au-Pt NRs show highly enhanced catalytic activity, which is due to the synergistic effect of bimetallic nanoparticles. The catalytic activity of immobilized Au-Pt NRs can be modulated by the volume transition of thermosensitive microgels. This demonstrates that core-shell microgels are capable of serving as “smart nanoreactors” for the catalytic active bimetallic nanoparticles with controlled morphology and high colloidal stability.

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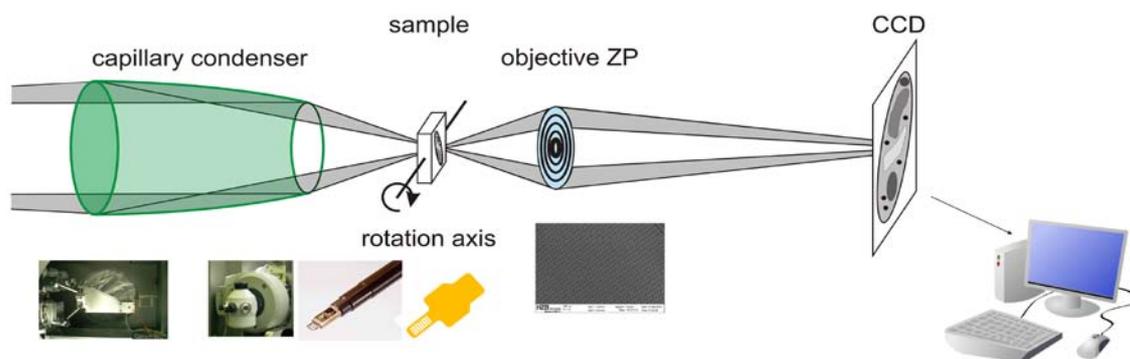
X-Ray Microscopy

Gerd Schneider

The research work in the Microscopy team comprises the development of advanced optical methods and instrumentation for nanoscale X-ray photonics. The group developed and operates one of the most advanced X-ray microscopes worldwide which permits NEXAFS spectromicroscopy on nanometer length scales with high energy resolution in the soft X-ray photon energy range. In addition, it is the first of its kind which permits high resolution tomography of mammalian cells using their native absorption contrast provided mainly by carbon and oxygen.

The Microscopy group is also one of the world's leaders in the development of high resolution diffractive X-ray optics. With the help of a state-of-the-art electron beam lithography system, special efforts are made towards sub-10 nm focusing. These experimental developments are complemented by theoretical studies based on rigorous coupled wave theory of the volume diffraction of high aspect ratio Fresnel zone plates. In-house developments include also new methods for advanced phase contrast methods combined with 3-D techniques.

Scientific applications comprise materials, environmental and life science. In materials science our main focus is the stress-induced migration and electromigration inside advanced integrated circuit structures. These structures are on the 100 nm length scale and need to be buried inside dielectrics. The X-ray microscope permits due to the high penetration depth of X-rays the imaging of much thicker samples than the TEM which opens up new experiments. These studies are performed in cooperation with industry and the Fraunhofer IZFP Dresden. In life science our scientific focus is the cell nucleus which is a true 3-D nanoscale structure. Cryo TEM gives us only images of thin sections whereas the cryo full-field X-ray microscope yields the complete cell including the nucleus with its three-dimensional structures like the nuclear membrane channels. In the futures, we want to develop methods for the correlation of fluorescence microscopy and 3-D tomography to provide a widely applicable technique to combine the biochemical information provided by labeled biomolecules in cells with structural information obtained by nano-tomography.



Coworkers:

PD Dr. Gerd Schneider
Dr. Peter Guttman
Dr. Stefan Rehbein
Dr. Stephan Werner
Katja Henzler

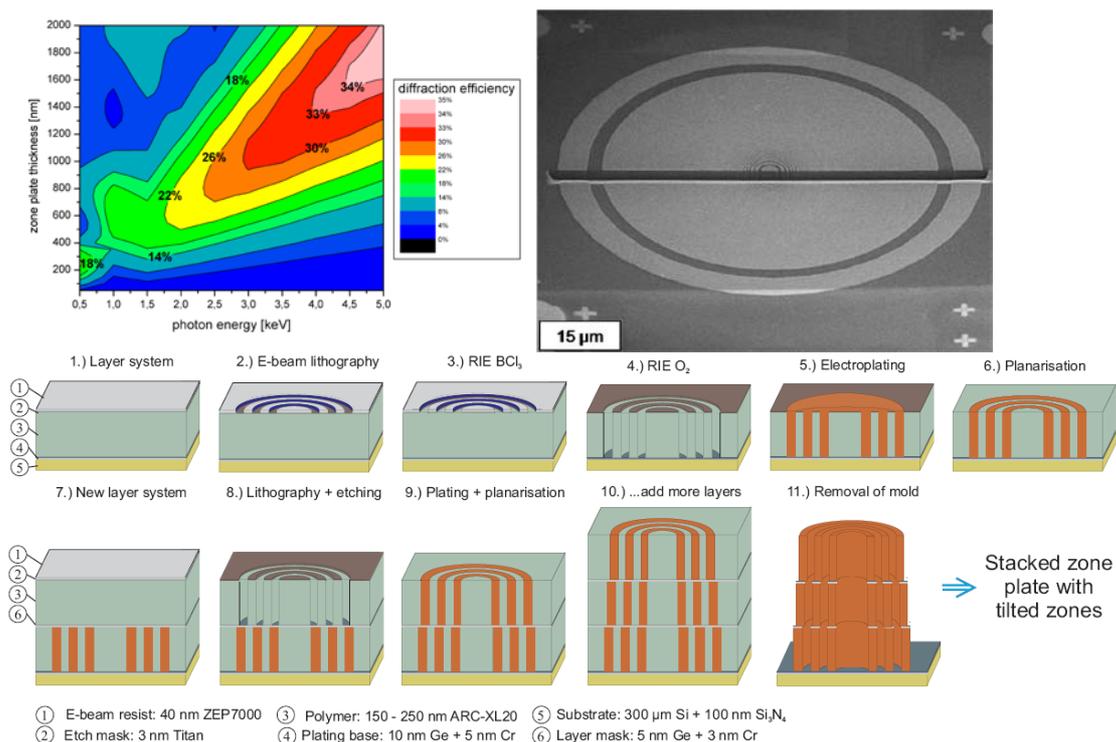
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Development of Diffractive Optics for High-Resolution X-Ray Imaging

S. Werner, S. Rehbein, G. Schneider

In the last decade advances in nanostructuring technology lead to rapid progress of the diffractive X-ray optics quality; they became the key elements for high resolution and energy resolving X-ray imaging techniques performed at synchrotron sources. The performance of Fresnel zone plates is characterized mainly by two parameters: The outermost zone width determines the numerical aperture and the height of the zone profile their diffraction efficiency. In the soft X-ray region the ratio of the zone height to the zone width is about 10:1. As sub-25 nm resolution optics are not commercially available, the microscopy group uses the VISTEC e-beam writer for pattern generation and in-house nanotechnology to manufacture these X-ray lenses. For manufacturing zone plates, different steps as thin layer technology, reactive ion etching and electroplating are required. Currently, the resolution obtainable with the HZB zone plates is about 10 nm. Electrodynamical calculations predict that in the future advanced stacked zone plates can combine high resolving power in the sub-10 nm range and high efficiency.

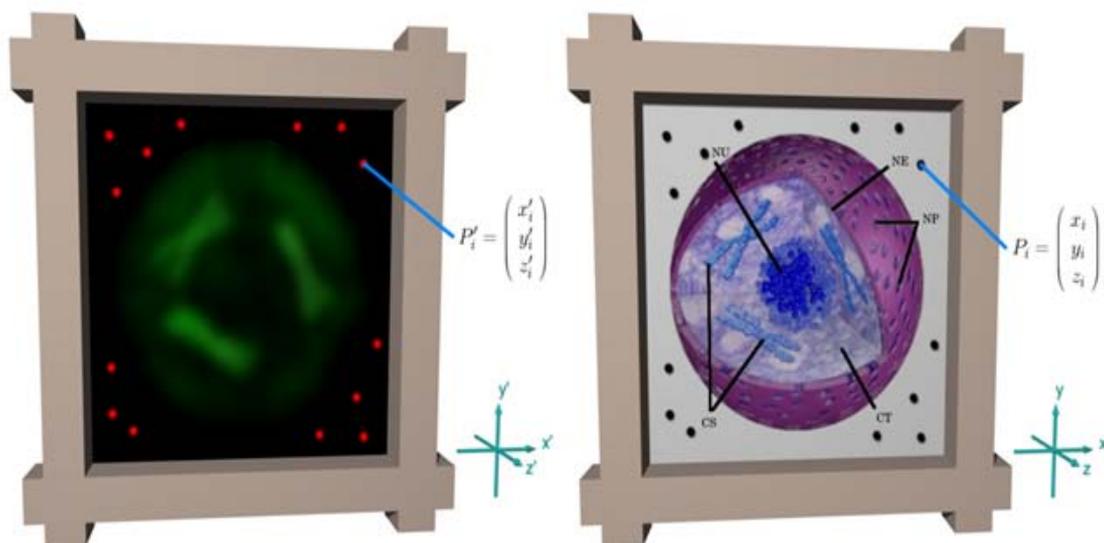


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Correlative 3-D Microscopy for Life Sciences

P. Guttman, G. Schneider

Fluorescence microscopy is an established technique in biophysical investigations of cells and cell nuclei, whereas 3-D X-ray microscopy is a relatively new approach with great potential which enables imaging of whole hydrated cells without chemical fixation, drying or slicing techniques as required in electron microscopy. Conventional optical fluorescence images are diffraction-limited to ~200 nm, whereas current X-ray images can achieve a ten-fold improvement in resolution. The interaction of X-rays is element specific; therefore, X-ray nano-tomography can be used to quantify the packing density of organic material. However, different proteins or molecular structures cannot be distinguished directly in X-ray microscope images. This problem is solved by the availability of specific fluorescent probes detectable by fluorescence microscopy. Thus the two imaging modalities are complementary. Since fluorescence and X-ray microscopy permit analysis of whole cells, it is possible to investigate the same cell in both microscopes.



The fluorescence microscope image (left) will provide information about the location of the labeled structures in the cell nucleus and the position (x'_i, y'_i, z'_i) of the markers (fluorescent dots outside the cell). The tomographic reconstruction obtained from the data acquired with the X-ray microscope reveals the internal nuclear structures (e.g., chromosomes, nucleolus, nuclear envelope, nuclear pores, and chromatin). These correlative studies are ideally suited for X-ray microscopy because of its ability to image whole cells in 3-D. This enables high throughput imaging of structures larger than a few hundred nanometer, which would otherwise be extremely time-consuming to locate and then serially reconstruct using correlative fluorescence and cryo electron microscopy of thin cell sections. With correlative microscopy, we expect to develop a widely applicable technique that, as applied to nuclear structure, will yield significant new insights.

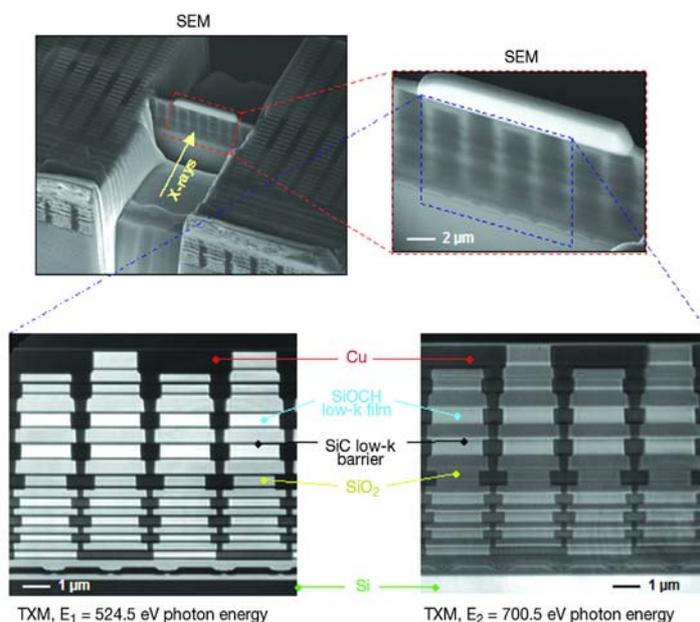
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Materials Science Applications

P. Guttman, S. Rehbein, S. Werner, G. Schneider, E. Zschech*

High-resolution X-ray imaging with a spatial resolution in the 10 - 30 nm range offers unique capabilities for process development and failure analysis in semiconductor industry. Buried metal interconnect structures like copper on-chip interconnects and through silicon vias (TSV) for 3D IC integration can be studied with excellent element-specific contrast. In addition, X-ray nano-tomography permits to study the kinetics of reliability-limiting processes like electromigration (EM) or stress-induced voiding (SIV).

X-ray microscopy is superior to SEM imaging if the structures are embedded as required for in-situ experiments. Such in-situ TXM experiments have been performed that give the ability to visualize mass transport processes and interconnect degradation while stressing fully embedded copper via/line test structures, applying accelerated test conditions (high temperature, high current density). These experiments allow to understand weaknesses in the interconnect technology that cause reliability-related failures. Such real-time imaging of interconnect degradation processes like electromigration and stress migration provide an ability to forecast the effect of process and materials changes on interconnect reliability and to optimize interconnect design rules. Particularly, void evolution in interconnects can be shown for Cu/low- κ structures with high spatial resolution, and rapid pathways for the directed mass transport and weak interfaces can be identified.



SEM micrograph (upper left) showing the FIB prepared fully passivated copper interconnect structures. Due to inelastic electron scattering in the passivation layer, the resolution in the magnified SEM micrograph (upper right) is limited. Images were taken at 525.5 eV and 700.5 eV energy using photons that traversed the prepared lamella in the X-ray microscope. Due to the different absorption properties of the used dielectric materials, these X-ray images reveal the different dielectric layers and the copper interconnects with a lateral resolution of about 20 nm.

dielectric layers and the copper interconnects with a lateral resolution of about 20 nm.

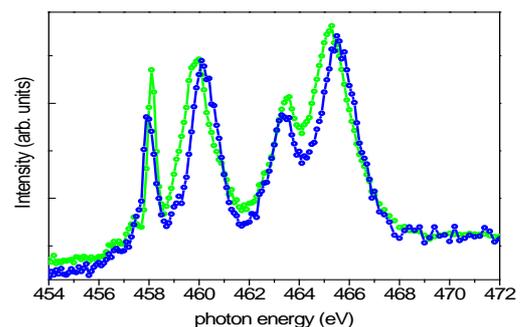
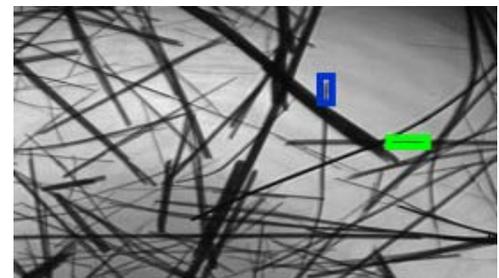
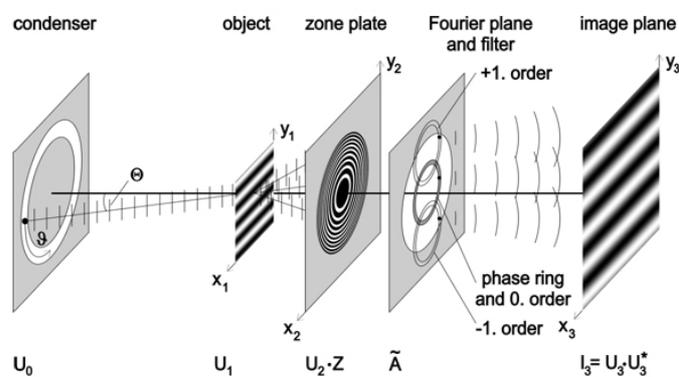
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X-ray Optical Methods and Instrumentation for Advanced X-ray Microscopy

G. Schneider, P. Guttman, S. Rehbein, S. Werner

In the nano-ages new tools for the analysis of complex structures is essential. So far the nano-world was mainly inspected by electron microscopy using a variety of different methods to utilize the image contrast formed by the interaction of electrons with matter. X-ray imaging methods are relatively new and much less developed compared to traditional microscopy techniques. However, they provide at least the same variety of interactions with matter to detect specific elements or chemical bonds. All this is based on high quality X-ray optics and advanced X-ray optical setups which take into account the relatively low efficiency of X-ray optics. Therefore, one goal of the X-ray microscope group is the development of novel methods for X-ray imaging to make use out of the unique interactions of X-rays with matter. X-ray optical setups providing a high energy resolution are required for spatially-resolved NEXAFS, special illumination schemes need to be developed for high-resolution phase contrast imaging or small focal spots using spatially coherent X-ray beams are required for nano-focusing. In addition, for the new instruments and applications special sample environments are required, for example cryo temperatures. Investigations of functionalized samples need linear or circular polarized light to visualize their inherent properties (e.g. polarization dependence of NEXAFS spectra of TiO_x nano-ribbons as shown in the figure).



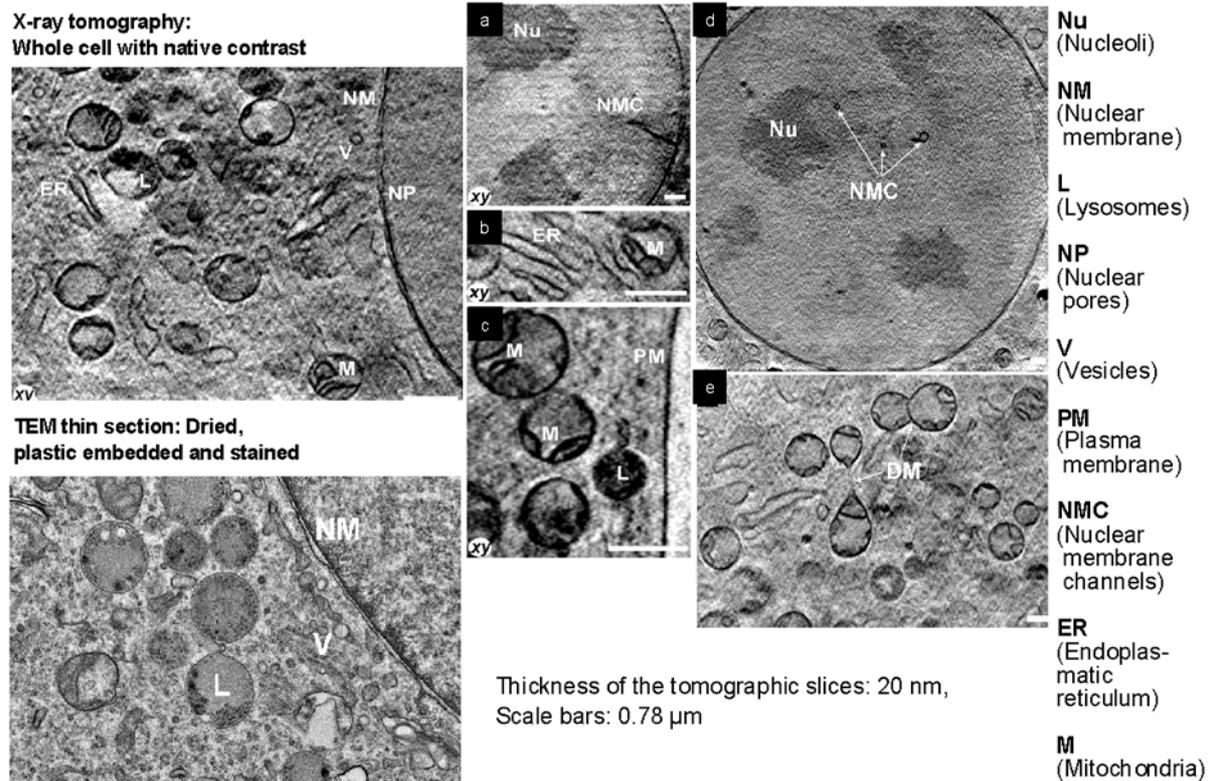
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Nano-Tomography of Cells

G. Schneider, P. Guttman, S. Werner, J. McNally*

Soft X-ray microscopy allows obtaining nano-scale 3D images of intact cells using only the natural contrast afforded by the different absorption or phase shift of organic matter and water. This permits entire cells to be examined while in their native state without chemical fixation, chemical staining or physical sectioning, but with only cryo-preservation.

Currently, the only alternative for visualizing 3D mammalian cell ultrastructure is cryo electron tomography performed on multiple serial sections, each of a thickness of much less than a micron, which must then be aligned to produce a 3D image. This is a painstaking process that requires 2-3 weeks per cell. Much faster nano-scale 3D imaging of intact cells can be performed with fluorescence super-resolution microscopy. However, this approach is fundamentally limited to examining the distribution of a few molecular markers per cell, and is therefore incapable of resolving a full spectrum of ultrastructural features. Thus X-ray microscopy with its potential to reveal the 3D ultrastructure in intact cells with a thickness of 10 μm fills an existing gap in current microscopy methods.



The new generation X-ray microscope at HZB allows routinely to visualize the plasma membrane, nuclear membrane, nuclear pores, nucleoli, endoplasmic reticulum, vesicles, lysosomes and mitochondria. It is now also possible to resolve internal organellar structures, such as mitochondrial cristae, the double nuclear membrane and lysosomal inclusions.

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* National Cancer Institute, NIH, Bethesda, USA

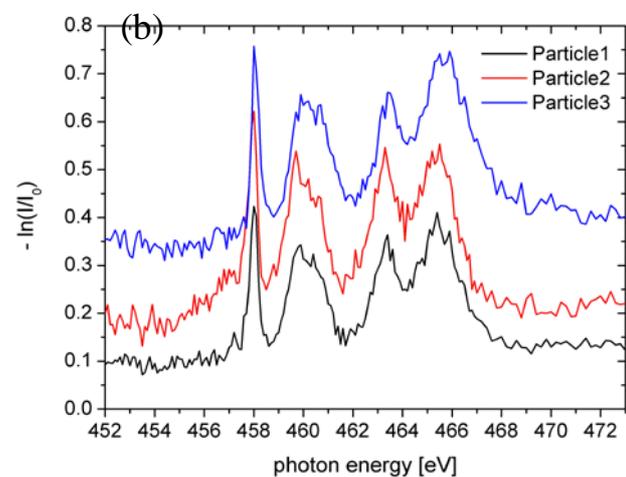
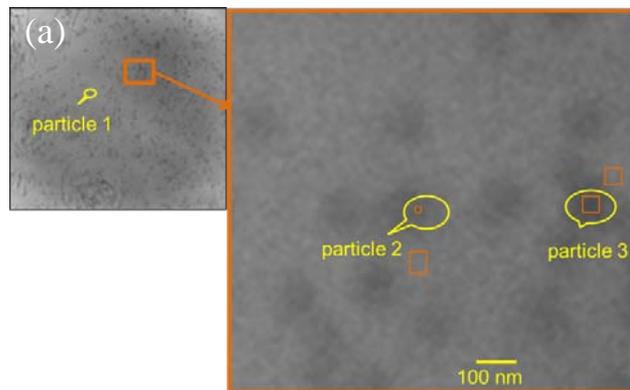
NEXAFS/cryo-TXM on colloidal hybrid structures

Peter Guttman, Katja Henzler, Yan Lu, Gerd Schneider

The properties of colloidal organic-inorganic hybrid materials are significantly affected by their morphology and elemental composition [1,2]. Therefore, suitable analytical methods are required for the investigation of these systems in situ. In order to investigate the morphology and composition of such particles in their solvated state, full-field cryo transmission X-ray microscopy (TXM) with automated nano-spectroscopy was used [3]. The investigated hybrid particles consist of a spherical polyelectrolyte brushes (SPB) which act as a carrier system for metal or metal oxide nanoparticles [1,2].

In the presented work we investigated titanium dioxide (TiO_2) nanoparticles on SPB by nano-spectroscopy at the Ti L-edge and O K-edge. The TiO_2 nanoparticles were synthesized by a modified sol-gel-process [1,2]. The diameter of these titanium dioxide particles was around 12 nm [1], which is in the resolution range of the HZB cryo-TXM at the BESSY II U41 undulator beamline [3]. The introduced method offers the possibility to distinguish between the different crystal structures of titanium dioxide. This is essential since only nanoparticle with an anatase crystal structure shows a high photolytic activity.

Figure (a) shows a TXM-image between 458 eV- 465.4 eV. Additionally, the used particles for the recording of the NEXAFS spectra at the Ti L-edge (Figure (b)) are highlighted. From the presented NEXAFS spectra in Figure (b) the electronic structure of the titanium dioxide can be calculated. Consequently, the crystal structure of titanium dioxide nanoparticles can be determined. The thorough evaluation of the presented spectra in Figure (b) shows that we have indeed an anatase crystal structure. This is in accordance with X-ray diffraction data we have collected on dried samples of this system.



(a) TXM-image of spherical polyelectrolyte brush with titanium dioxide nanoparticles. Highlighted are the particles which were used to record the NEXAFS spectra (b) at the Ti L-edge.

(b) NEXAFS spectra at the Ti L-edge of titanium oxide nanoparticles. The careful analysis of the recorded spectra reveals that the TiO_2 nanoparticles have an anatase crystal structure.

- [1] Y. Lu, M. Hoffmann, R.S. Yelamanchili, A. Terrenoire, M. Schrunner, M. Drechsler, M.W. Möller, J. Brey, M. Ballauff, *Macromol. Chem. Phys.* **2009**, 210 (5), 377 – 386.
 [2] Y. Lu, T. Lunkenbein, J. Preussner, S. Proch, J. Brey, R. Kempe, M. Ballauff, *Langmuir* **2010**, 26 (6), 4176-4183.
 [3] S. Heim, P. Guttman, S. Rehbein, S. Werner, G. Schneider, *J. Phys. Conf. Series* **2009**, 186, 0122041.

Soft Matter Theory Group: Methods, Mission, and Research

Joachim Dzubiella

The newly formed theory group establishes another important pillar of the Soft Matter and Functional Materials Institute at the HZB by providing *theoretical models* of soft (biological) condensed matter systems using *analytical calculations and computer simulations*. The topics studied by the group cover a broad range of systems and include, for instance, the equilibrium and nonequilibrium structure and phase behavior of colloidal and polymeric fluids, protein structure and dynamics in aqueous (poly)electrolyte solutions, or self-organization of soft organic materials at interfaces.

Inherent to any complex soft matter system is the appearance of multiple length and time scales. In order to tackle the multiscale challenge theoretically, the many small (and fast) degrees of freedom need to be integrated out, what the theorists call 'coarse-graining'. The latter can be achieved by combining various theoretical treatments. Here, the theory group employs approaches from classical statistical mechanics, liquid state theory, and various computer simulation techniques. In particular, the group performs Molecular dynamics (MD), Brownian dynamics (BD), or Monte-Carlo (MC) simulations combined with integral equation theory or (classical) dynamical density functional theory.

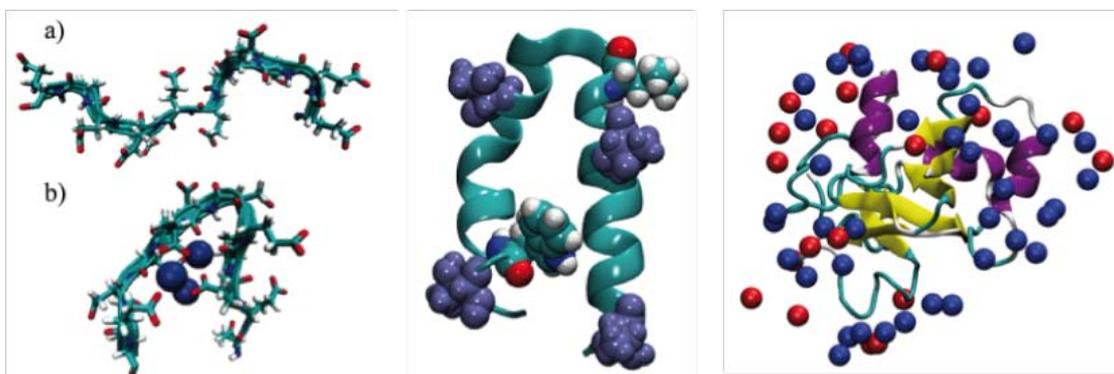
The computationally demanding MD simulations of molecular systems will be mainly performed on the high performance cluster Dirac directly located and maintained on the Lise-Meitner-Campus. The cluster currently features 20 nodes embracing 256 computing processor units (CPUs) and parallel batch processing software and state-of-the-art molecular dynamics simulation software. Given the excellent personal and spatial infrastructure at the HZB the cluster will be extended by another 144 processor units in near future which warrants competitive computer power for the planned theoretical undertakings at the Soft Matter Institute.

The theory group thus directly complements the experimental efforts in soft matter physics at the HZB. It provides deeper physical insights by modeling and boiling down the complex reality to simple model systems. Atomistically-resolved MD simulations on the other hand enlighten the molecular mechanisms behind certain phenomena inaccessible for direct measurement or spatial resolution. Last but not least, the theory group will support and guide software developments for analyzing and modeling structural data obtained from HZB beamline users. The strong synergy between the theory and experiments concentrated here at the HZB will hopefully result in fascinating discoveries in soft material science in the coming years.

Examples of planned projects:

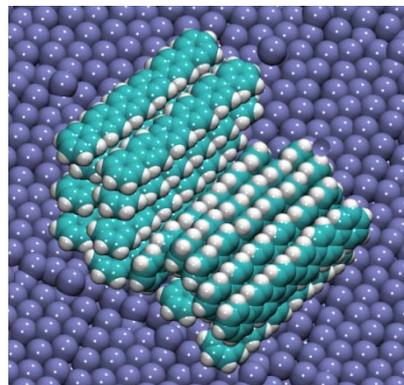
One planned direction of the theoretical research is the rheology of driven colloids, e.g., by shear or a gravitational field. Using BD computer simulations and dynamic density functional theory (DDFT) nonequilibrium structural evolution in time can be theoretically predicted [1]. Future research will extend these investigations to more complicated particle geometries (e.g., dumbbells) and other external fields, e.g., shear as performed experimentally at the colloidal lab of the Soft Matter Institute at the HZB.

Another planned direction is the investigation of ionic and hydration effects on protein interactions, protein dynamics and function. It is well established that solvation of proteins by water and specific ions has fundamental effects on protein folding and function and is strongly coupled to the protein dynamical modes. The theory group has strong expertise on modeling solvation effects by atomistically resolved MD computer simulations on the folding of oligomers [2,3], dynamical relaxation of peptidic polyelectrolytes [4], or ions binding to proteins. Further research will be tightly connected to the time-resolved neutron scattering experiments at the HZB to probe protein dynamics and its relation to protein function.



MD simulation snapshots of biomolecules. Left: polyglutamic acid a) extended, b) collapsed with counterions [4]. Middle: kinked anti-freeze protein with water-active threonine side chains (blue) [3]. Right: halophilic (salt-loving) ferredoxin with surface-bound ions.

The theory is group is also involved in a long-term enterprise to explore the morphology of hybrid inorganic-organic systems (HIOS) within a planned Berlin-wide Collaborative Research Center. The aim of the theory project is to support experimental microscopy and spectroscopy measurements by atomistically-resolved molecular dynamics computer simulations. The fundamental molecular insight will help interpreting and guiding the experimental efforts with regard to structure nucleation and growth of adjunct organic molecules close to inorganic surfaces. This will include, for instance, p-sexiphenyl molecules adjacent to ZnO, as shown in the preliminary simulation snapshot on the right.



- [1] Nonequilibrium sedimentation of colloids on the particle scale
P. Royall, J. Dzubiella, M. Schmidt, and A. van Blaaderen, *Phys. Rev. Lett.* 98, 188304 (2007)
- [2] Salt-specific stability and denaturation of a short salt-bridge forming alpha-helix
J. Dzubiella, *J. Am. Chem. Soc.* 130(42), 14000-14007 (2008).
- [3] The A17L mutation of wf-AFP1 reduces anti-freeze activity by kinking the helix and reducing long-range protein-water coupling, S. Ebbinghaus, J. Dzubiella, et al. *Biophys. Lett.*, submitted (2010)
- [4] Molecular insights into the ion-specific kinetics of anionic peptides
J. Dzubiella, *J. Phys. Chem. B* 114, 7098 (2010).

Polymer Physics

Dr. Sebastian Seiffert

Polymer gels consist of a three dimensional network of crosslinked polymer chains which are swollen in a solvent; they are fascinating materials for applications as superabsorbers, matrixes in analytical chemistry or biology, or as storage and delivery systems for actives. Classically, network junctions are formed by chemical bonds, ensuring great mechanical stability. However, these chemical crosslinks are permanent, such that chemical gels cannot be processed or recycled. Permanent chain interconnection is also detrimental for encapsulation and controlled release applications. It is therefore desirable to use *reversible* gels, which is readily achieved by *supramolecular* chain crosslinking. Previous work in this field has produced various types of such materials, primarily realized through chain interconnection by hydrogen bonding or transition metal complexation. However, a comprehensive characterization and deep understanding between the chemical structure and the phenomenological behavior of these promising materials is still lacking.

Our research focuses on polymer networks that are crosslinked by supramolecular bonds. We prepare, study, and process these networks in a systematic fashion. For this purpose, we use a universal covalent precursor polymer and equip it with side groups that can be interconnected by non-covalent interactions such as hydrogen bonding or transition metal complexation. This leads to networks that consist of the same basis material, yet exhibiting a strongly varying strength of chain interconnection.

We study these supramolecular gels to derive fundamental relations between the strength of non-covalent crosslinking and the network structure and dynamics, using methods such as macroscopic rheology as well as light, neutron, and x-ray scattering. In addition, we use heterophase techniques such as miniemulsification and droplet-based microfluidics to fabricate supramolecular nano- and microgel particles. These particles can serve as nano- and microcapsules for the encapsulation and controlled release of actives, including drugs, biopolymers, and living cells. We also use these particles as microscopic probes to study their polymer network architectures, and we investigate the physical chemical properties of densely packed suspensions of these micro- and nanogels.



Seiffert Group
FU Berlin – Dpt. of Chemistry
HZB – Soft Matter & Funct. Mater.

- Physical Chemistry of Supramolecular Gels
- Microfluidic Formulation of Supramol. Microgels
- Structure and Dynamics of Microgel Suspensions

Coworkers:

Dr. Sebastian Seiffert

M.Sc. Torsten Rossow

Publications in 2008–2011:

S. Seiffert, J. Dubbert, W. Richtering, D. A. Weitz, "Reduced UV light scattering in PDMS microfluidic devices." *Lab Chip* **2011**, *11*, 966–968.

D. Steinhilber, S. Seiffert, J. A. Heyman, F. Paulus, D. A. Weitz und R. Haag, "Hyperbranched polyglycerols on the nanometer and micrometer scale." *Biomaterials* **2011**, *32*, 1311–1316.

S. Seiffert, D. A. Weitz, "Microfluidic fabrication of smart microgels from macromolecular precursors." *Polymer* **2010**, *51*, 5883–5889.

S. Seiffert, M. B. Romanowsky, and D. A. Weitz, "Janus Microgels Produced from Functional Precursor Polymers." *Langmuir* **2010**, *26*, 14842–14847.

S. Seiffert and D. A. Weitz, "Controlled Fabrication of Polymer Microgels by Polymer-Analogous Gelation in Droplet Microfluidics." *Soft Matter* **2010**, *6*, 3184–3190.

S. Seiffert, J. Thiele, A. R. Abate, and D. A. Weitz, "Smart Microgel Capsules from Macromolecular Precursors." *J. Am. Chem. Soc.* **2010**, *132*, 6606–6609.

S. Seiffert and W. Oppermann, "Diffusion of Linear Macromolecules and Spherical Particles in Semidilute Polymer Solutions and Polymer Networks." *Polymer* **2008**, *49*, 4115–4126.

Soft Matter and Functional Materials

CVs



Prof. Dr. Matthias Ballauff

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Email: matthias.ballauff@helmholtz-berlin.de



Area of Expertise and Field of Interest

Area of Expertise: Physics of colloids and macromolecules

Field of Interest: Chemistry and physics of colloidal systems, scattering methods

Scientific Development

Study	1971- 1977 Study of chemistry at the University of Mainz
PhD	1981 at the University of Mainz (Prof. Dr. B. A. Wolf)
Post-doc	1981-1983 Dept. Chemistry, Stanford University, Prof. Dr. P. J. Flory
	1994- 1999 Research Associate, Max-Planck-Institut für Polymerforschung, Mainz
	1999 Habilitation at University of Mainz
Full Chair	1990- 2003 at University of Karlsruhe
	2003- 2009 at University of Bayreuth
	Since 2009 Head of the Institute for Soft Matter and Functional Materials, HZB, Berlin, and Professor of Physics, Humboldt Universität zu Berlin

Selected Scientific Expert, Review or Council Activities, Awards

Member of the DFG-Panel for the special research areas of the Deutsche
Forschungsgemeinschaft (Mitglied des DFG Ausschusses für die Angelegenheiten der
Sonderforschungsbereiche) (1999-2005), Themenkommission Deutsche Bunsen-
Gesellschaft (2008 – 2011), Member of „Ständiger Ausschuss“ Deutsche Bunsen-
Gesellschaft (2011 – 2012), Editor Polymer (since 2004), Scientific Advisory Committee
HERCULES

Invited Talks (selection)

“8th International Conference on Advanced Polymers via Macromolecular Engineering”,
Dresden 2009; “44th Biennial Meeting of the German Colloid Society”, Hamburg 2009; “43rd
IUPAC World Polymer Congress Macro”, Glasgow, England 2010; “23 General Conference
of the European Physical Society”, Warschau, Polen 2010; “RUSNANOTECH Forum”,
Moskau, Russland 2010; “8th International Symposium on Polyelectrolytes”, Shanghai, China
2010

Selected publications

1. J. M. Brader, M. Siebenbürger, M. Ballauff, K. Reinheimer, M. Wilhelm, S. J. Frey, F. Weysser, M. Fuchs, *Nonlinear response of dense colloidal suspensions under oscillatory shear: Mode-coupling theory and Fourier transform rheology experiments*, Phys. Rev. E 2010, **82**, 061401
2. M. Hoffmann, M. Siebenburger, L. Harnau, M. Hund, C. Hanske, Y. Lu, C. S. Wagner, M. Drechsler, M. Ballauff, *Thermoresponsive colloidal molecules*, Soft Matter, 2010, **6**, 1125-1128
3. K. Henzler, B. Haupt, K. Lauterbach, A. Wittemann, O. Borisov, M. Ballauff, *Adsorption of beta-Lactoglobulin on Spherical Polyelectrolyte Brushes: Direct Proof of Counterion Release by Isothermal Titration Calorimetry*, J Am Chem Soc, 2010, **132**, 3159-3163
4. H. H. Winter, M. Siebenbürger, D. Hajnal, O. Henrich, M. Fuchs, M. Ballauff, *An empirical constitutive law for concentrated colloidal suspensions in the approach of the glass transition*, Rheol Acta, 2009, **48**, 747-753
5. J. J. Crassous, C. N. Rochette, A. Wittemann, M. Schrinner, M. Ballauff, M. Drechsler, *Quantitative Analysis of Polymer Colloids by Cryo-Transmission Electron Microscopy*, Langmuir **25**, (2009) 7862.
6. M. Schrinner, M. Ballauff, Y. Talmon, Y. Kauffmann, J. Thun, M. Möller, J. Breu, *Single-Nanocrystals of Platinum Prepared by Partial Dissolution of Au-Pt Nanoalloys*, Science, **323**, (2009) 617.
7. K. Henzler, S. Rosenfeldt, A. Wittemann, L. Harnau, S. Finet, T. Narayanan, M. Ballauff, *Directed motion of proteins along tethered polyelectrolytes*, Phys. Rev. Lett., **100**, (2008) 158301.
8. M. Ballauff, *Spherical Polyelectrolyte Brushes*, Progr. Polym. Sci., **32**, (2007) 1135.
9. Y. Lu, Y. Mei, M. Drechsler, M. Ballauff, *Thermosensitive Core-Shell Particles for Ag-Nanoparticles: Modulating the Catalytic Activity by the Volume Transition in Networks*, Angew. Chemie Intl. Ed. **45**, (2006) 813.

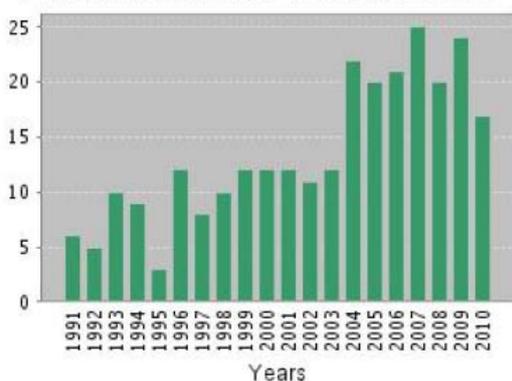
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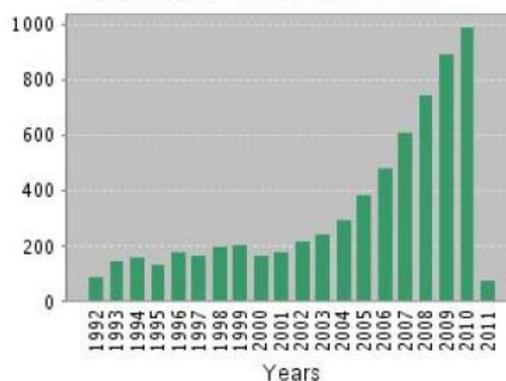
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Area of Expertise and Field of Interest

Area of Expertise: Neutron and X-ray instrumentation

Field of Interest: Physics of colloidal systems, Deposition and characterization of multilayer systems

Scientific Development

Study 1981- 1989 Study of physics at the Technische Universität Berlin
PhD 1993 at the Technische Universität Berlin (Prof. Dr. F. Mezei)
Post-doc 1994-2001 Laboratory for Neutron Scattering Villigen, Paul Scherrer
Institut and ETH Zürich, Prof. Dr. A. Furrer / Prof. Dr. P. Böni
Staff positions 2001- 2002 Staff scientist, Laboratory for Neutron Scattering Villigen,
Paul Scherrer Institut and ETH Zürich
Since 2002 Staff scientist, Dept. Methods and Instrumentation for
Neutron Scattering and Institute for Soft Matter and Functional
Materials, HZ Berlin

Selected Scientific Expert, Review or Council Activities, Awards

International instrument advisory team for the QUOKKA-SANS, Australian Nuclear Science and Technology Organisation (until 2005), Advisory committee of Workshop on Neutron Delivery Systems, ILL Grenoble (1.-3.7.'09), International advisory committee of the Laboratoire Léon Brillouin of CEA/CNRS, Saclay (since 2010)

Invited Talks (selection)

KAERI Seminar, Daejeon, 2006, Seminar of the Laboratoire Léon Brillouin of CEA/CNRS, Saclay, 2006

Selected publications

- [1] Y. Gerelli, M.T. Di Bari, A. Deriu, D. Clemens, L. Almásy, *Lipid multilayered particles: the role of chitosan on structure and morphology*, *Soft Matter* **6**, 2533–2538 (2010)
- [2] F. Cousin, J. Gummel, D. Clemens, I. Grillo, F. Boué, *Multiple Scale Reorganization of Electrostatic Complexes of Poly(styrenesulfonate) and Lysozyme*, *Langmuir*, **26**, 7078–7085 (2010)
- [3] M.G. Ortore, R. Sinibaldi, F. Spinozzi, F. Carsughi, D. Clemens, A. Bonincontro, P. Mariani, *New Insights into Urea Action on Proteins: A SANS Study of the Lysozyme Case*, *J. Phys. Chem. B* **112**, 12881-12887 (2008).
- [4] I. Estrela-Lopis, S. Leporatti, E. Typlt, D. Clemens, E. Donath, *Small Angle Neutron Scattering Investigations (SANS) of Polyelectrolyte Multilayer Capsules Templated on Human Red Blood Cells*, *Langmuir*, **23**, 7209–7215 (2007).
- [5] V.-M. Graubner, D. Clemens, Th. Gutberlet, R. Kötz, Th. Lippert, O. Nuyken, B. Schnyder, A. Wokaun, *Neutron Reflectometry and Spectroscopic Ellipsometry Studies of Cross-Linked Poly(dimethylsiloxane) after Irradiation at 172 nm*, *Langmuir* **21**, 813 (2005).

Prof. Dr. Joachim Dzubiella

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Area of Expertise and Field of Interest

Area of Expertise: Theoretical Physics of Complex Fluids

Field of Interest: Colloidal systems, Aqueous electrolytes solutions, Proteins

Scientific Development

- 1994-1999 Study of Physics at the HH-University of Düsseldorf
- 2002 PhD at the HH-University of Düsseldorf (Prof. Dr. H. Löwen)
- 2002-2004 Postdoc Dept. Chemistry, Cambridge University, UK, Prof. Dr. J.-P. Hansen
- 2004-2006 Postdoc Dept. Biochemistry, Fellow of the Center for Theoretical Biophysics, UC San Diego, USA (Prof. J. A. McCammon).
- 2006-2010 Emmy-Noether Fellow, Research Group Leader, Physics Dept., Technical University Munich
- Since 2010 Group Leader at the Soft Matter and Functional Materials Institute, HZB, Berlin, and Professor of Physics, Humboldt Universität zu Berlin

Scholarships

- 2002 DAAD Auslandsstipendium (UC Santa Barbara)
- 2004 DFG Forschungsstipendium (UC San Diego)
- 2006 DFG Emmy-Noether Fellowship (TU Munich)

Invited Talks (selection)

2007, UTAM Symposium on Swelling and Shrinking of Porous Materials: From Colloid Science to Poromechanics, Rio de Janeiro, Brazil; 2008, CTBP summer school "Coarse-Grained Physical Modeling of Biological Systems: Advanced Theory and Methods"; UCSD, USA; 2009, International Workshop on Continuum Modeling of Biomolecules, Beijing, China; 2010 Gordon, Research Conference on "Aqueous Systems", USA; 2010 Pacificchem2010, Honolulu, USA

Selected publications

- 1) Ionic-specific excluded-volume correlations and solvation forces
I. Kalcher, J. C. F. Schulz, and J. Dzubiella, *Phys. Rev. Lett.* **104**, 097802 (2010).
- 2) Dewetting-controlled binding of ligands to hydrophobic pockets
P. Setny, Z. Wang, L.-T. Cheng, B. Li, J. A. McCammon, and J. Dzubiella, *Phys. Rev. Lett.* **103**, 187801 (2009).
- 3) Sequence-specific size, structure, and stability of tight protein knots
J. Dzubiella, *Biophys. J.* **96**, 831-839 (2009).
- 4) Salt-specific stability and denaturation of a short salt-bridge forming alpha-helix
J. Dzubiella, *J. Am. Chem. Soc.* **130**(42), 14000-14007 (2008).
- 5) Nonequilibrium sedimentation of colloids on the particle scale, . Royall, J. Dzubiella, M. Schmidt, and A. van Blaaderen, *Phys. Rev. Lett.* **98**, 188304 (2007).

Dr. Günter Johannes Goerigk

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Area of Expertise and Fields of Interest

Area of Expertise: Anomalous Small-Angle X-ray Scattering, Very Small-Angle Neutron Scattering, Decomposition kinetics in alloys

Fields of Interest: Time resolved in-situ ASAXS studies on the decomposition kinetics of Copper-Cobalt alloys. Extensive investigations of amorphous Silicon-Germanium alloys used in solar cell techniques. Analysis of conformational and quantitative properties of counter ion condensation to polyelectrolytes by anomalous small-angle X-ray scattering. Critical scattering of amorphous ternary metallic alloys. Critical Casimir effect in binary liquid mixtures analyzed by V-SANS (Very Small-Angle Neutron Scattering). Numerous ASAXS studies on alloys, ceramics, magnetic systems, catalysts, semiconductors, glasses, polymers, membranes and soft matter systems in collaboration with scientific groups from Europe, United States, South America and Asia.

Scientific Development

Study	1973 - 1982	Study of physics, Dipl.-Phys. at University of Bonn
PhD	1988	in chemistry at University of Hamburg (Prof. Dr. H. G. Zachmann, Prof. Dr. H.B. Stuhmann)
Post-doc	1988 -1990	GKKS Research Centre, Institute of Materials Research
Synchrotron radiation	1983 -1984	ASAXS beam line X15 at EMBL Outstation at DESY Hamburg, Prof. Dr. H.B. Stuhmann
	1985 - 1990	Establishing ASAXS beam line ROEFO1 together with B. Munk at HASYLAB/DESY Hamburg
	1991 - 2007	Scientist in charge at ASAXS beam line JUSIFA @HASYLAB/DESY Hamburg, Forschungszentrum Jülich
Neutrons	2007 -2010	Scientist at V-SANS beam line KWS-3@JCNS/FRM II Munich, Forschungszentrum Jülich, Comprehensive upgrade activities, commissioning, implementation into the regular FRM II proposal and scheduling system since December 2009
	since 10/2010	Institute of Soft Matter and Functional Materials, Helmholtz-Zentrum Berlin

Selected Scientific Expert, Review or Council Activities

1996-2005	Member of the scientific committee of DESY
2003-2007	Project leader JUSIFA@HASYLAB
2007-2010	Member of the co-ordination panel JCNS instrumentation at FRM II ASAXS-consultant

Invited Talks (selection)

Goerigk, G.; Williamson, D.L.; Huber, K.; Schweins, R.: Anomalous Small-Angle X-ray Scattering: A Precise Quantitative Method in Solid State Physics, Chemistry and Materials Science, Invited talk at the XIII International Conference on Small-Angle Scattering : Kyoto International Conference Hall, Kyoto, Japan, 09.07.2006 - 13.07.2006

Goerigk, G.: Anomalous Small-Angle X-ray Scattering: A Precise Quantitative Method in Solid State Physics, Chemistry and Materials Science, Invited talk at the SR User Meeting : Diamond Light Source, Didcot, Great Britain, 13.09.2007 - 14.09.2007

Goerigk, G.: Anomalous Small-Angle X-ray Scattering: A precise Quantitative Method for the Structural Analysis in Chemistry, Solid State Physics and Membrane Science – A view on new results and perspectives, 1st International ASAXS-Workshop, Berlin, Germany, 14.05.2009

Goerigk, G., Anomalous Small-Angle X-ray Scattering: A Precise Quantitative Method in Chemistry and Solid State Physics, Hungarian Academy of Science, MTA Kémiai Kutatóközpont, Budapest, Hungary, 30.06.2009

Goerigk, G., Anomalous Small-Angle X-ray Scattering: A Precise Quantitative Method in Condensed Matter Research, Invited talk to be held at REXS 2011, International Conference/School on Resonant Elastic X-Ray Scattering in Condensed Matter, Aussois, France; 13.-17. June 2011

Selected publications

Bota, A.; Varga, Z.; Goerigk, G.: Biological systems as nanoreactors: Anomalous small-angle X-ray scattering study of the CdS nanoparticle formation in multi-lamellar vesicles. *Journal of Physical Chemistry B* (2007), 111(8), 1911-1915.

Goerigk, G.; Huber, K.; Schweins, R. (2007): Probing the extent of the Sr²⁺ ion condensation to anionic polyacrylate coils: A quantitative anomalous small-angle x-ray scattering study. *Journal of Chemical Physics*, 127(15), 154908/1-154908/8.

Goerigk, G.; Mattern, N. Critical scattering of Ni-Nb-Y metallic glasses probed by quantitative anomalous small-angle x-ray scattering. *Acta Materialia* (2009), 57(12), 3652-3661.

Goerigk, G., Varga, Z., Comprehensive upgrade of the high-resolution small-angle neutron scattering instrument KWS-3 at FRM II, *Journal of Applied Crystallography* (2011), in press.

Lectures at University Paderborn, Department of Chemistry and Humboldt University Berlin, Institute of Physics

In summer semester 2010 in Physical Chemistry contributing with 8 hours to the lecture 'Structure determination' of Prof. K. Huber.

In addition to the lecture 'Structure determination' of summer semester 2010 practical exercises with 4 students of Universität Paderborn at Jülich Centre of Neutron Science (JCNS) at the Research reactor FRM II@TUM in Garching/Munich. Very small-angle neutron scattering experiments at V-SANS beamline KWS-3. Three days exercises, seminar and tutorials.

In winter semester 2010/2011 lectures and exercises in the lecture 'Polymer science' of Prof. J. Rabe and Prof. M. Ballauff at Humboldt Universität zu Berlin, Institut für Physik

Scientific co-operations (selected)

Prof. K. Huber, Universität Paderborn, Fakultät für Naturwissenschaften, Department Chemie, Warburgerstr. 100, D-33098 Paderborn, Federal Republic of Germany, Combined SANS, ASAXS and LS Studies of Structural Transformation in Polyacrylate Anions Induced by Specifically Interacting Metal Cations.

Dr. A. Bóta, Department of Biological Nanochemistry, Chemical Research Center, Hungarian Academy of Sciences, Pusztaszeri ut 59-67, H-1025, Budapest, Hungary, Structure and interparticle interactions in biomedical colloidal systems

Dr. Nikoline Hansen

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Area of Expertise and Field of Interest

Area of Expertise: Print and New Media, Assistance
Field of Interest: Organisation and Consulting

Scientific Development

5/1986	Magister in American Studies, Politics as well as Prehistoric and Protohistoric Archaeology
1/1996	PhD at the Freie Universität Berlin, American Studies Public Relations, DAPR New Media, Fachhochschule für Technik und Wirtschaft
06/1986-05/2009	Team Secretary at BESSY and BESSY II Project, Personal Assistant to Professor Jaeschke (Director, accelerator physics)
09/1997-10/1998	Night School Public Relations, DAPR Examination PR Consultant
11/2001-03/2003	Advanced off-the-job training in Visual Computing and Web Technology (Medieninformatik) at the Fachhochschule für Technik und Wirtschaft Berlin
Since 06/2009	Institute Soft Matter and Functional Materials, administrative officer, Helmholtz-Zentrum Berlin, Germany, Prof. Dr. M. Ballauff

Print Media:

BESSY II – Eine optimierte Undulator/Wiggler-Speicherring Lichtquelle für den VUV- und
XUV-Spektralbereich (1986)

Visions of Science: the BESSY SASE-FEL in Berlin-Adlershof Scientific Case for an FEL
(2001)

Technical Design Report BESSY FEL (2004), BESSY, Berlin

International Conference Organisation

FEL06 (Local Organisation Committee and Conference Secretary Berlin 2006)

Organisation of International Committees and Workshops

BESSY II MAC (1993-1998, 11 Meetings)

Workshop on Scientific Case for the BESSY SASE-FEL, Holzau 2001

Dr. Thomas Hauß

(1958)

F-I2 Soft Matter and Functional Materials,
Helmholtz-Zentrum Berlin für Materialien und
Energie GmbH, Hahn-Meitner-Platz 1, 14109 Berlin

Tel: 030 8062-42071 (-42999, Fax)

Email: hauss@helmholtz-berlin.de



Area of Expertise and Field of Interest

Area of Expertise: Biophysics

Field of Interest: Membrane biophysics, peptide-membrane interactions, protein dynamics, neutron scattering methods

Scientific Development

Study 1977- 1986 Study of physics, Technische Universität Berlin

PhD 1992 Freie Universität Berlin (Prof. Dr. M.P. Heyn)

Post-doc 1993-1996 Hahn-Meitner-Institut Berlin, Prof. Dr. M. Steiner

1996-1999 Research Scientist, Clemens-Schöpf-Institute, Technische Universität Darmstadt, Prof. Dr. N.A. Dencher

2000-2004 Research Scientist, Dept. Physics, Heinrich-Heine Universität Düsseldorf, Prof. Dr. G. Büldt

2005-2009 Research Scientist, Clemens-Schöpf-Institute, Technische Universität Darmstadt, Prof. Dr. N.A. Dencher

permanent Since 2009 Group leader Biophysics, Institute Soft Matter and Functional Materials, Helmholtz-Zentrum Berlin für Materialien und Energie

Selected Scientific Expert, Review or Council Activities, Awards

Reviewer for: Biotechnology and Biological Sciences Research Council (BBSRC), UK; NIST Center for Neutron Research, Gaithersburg, USA; DLab, ILL Grenoble, France.

Invited Talks (selection)

Workshop on (Glyco)lipids, structures, functions, and interactions, Universität Hamburg, Germany 2010; Neutrons in Biology, Lund University Sweden 2009; 13th International Conference on Retinal Proteins, Barcelona, Spain 2008; 3rd Japanese-French Seminar on Protein Dynamics, Grenoble, France 2007; 12th International Conference on Retinal Proteins - Satellite Meeting Nagoya, Japan 2006

Selected publications

- [1] A. Buchsteiner, T. Hauß, S. Dante, N. A. Dencher. *Alzheimer's disease amyloid-beta peptide analogue alters the ps-dynamics of phospholipid membranes*. *Biochimica et Biophysica Acta* **1798** (2010), 1969-1976
- [2] A. Schröter, D. Kessner, M. A. Kiselev, T. Hauß, S. Dante, R. H. H. Neubert. *Basic nanostructure of stratum corneum lipid matrices based on ceramides [EOS] and [AP]. A neutron diffraction study*. *Biophysical Journal* **97** (2009) 1104-14.
- [3] H. Seelert, D. N. Dani, S. Dante, T. Hauß, F. Krause, E. Schafer, M. Frenzel, A. Poetsch, S. Rexroth, H. J. Schwaßmann, T. Suhai, J. Vonck, N. A. Dencher. *From protons to OXPHOS supercomplexes and Alzheimer's disease: Structure-dynamics-function relationships of energy-transducing membranes*. *Biochimica et Biophysica Acta* **1787**, (2009) 657-671.
- [4] J. Pieper, A. Buchsteiner, N. A. Dencher, R. E. Lechner, T. Hauß. *Transient protein softening during the working cycle of a molecular machine*. *Physical Review Letters* **100**, (2008) 228103
- [5] S. Dante, T. Hauß, A. Brand, N. A. Dencher. *Membrane fusogenic activity of the Alzheimer's peptide A β (1-42) demonstrated by small-angle neutron scattering*. *Journal of Molecular Biology* **376**, (2008) 393-404.
- [6] S. Dante, T. Hauß, N. A. Dencher. 2006. *Cholesterol inhibits the insertion of the Alzheimer's peptide A β (25-35) in lipid bilayers*. *European Biophysics Journal* **35**, (2006) 523-531

Dr. Martin Hoffmann

(1981)

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Area of Expertise and Field of Interest

Area of Expertise: Polymer- and Colloid Chemistry

Field of Interest: Anisotropic colloids, polyelectrolytes, microgels, heterogeneous catalysis, depolarized dynamic light scattering

Scientific Development

10/2002 – 08/2007 Study of Polymer- and Colloid Chemistry at the University of Bayreuth

10/2007 – 08/2010 PhD at the University of Bayreuth (Prof. Dr. M. Ballauff):

“Synthesis and Characterization of Anisotropic Colloidal Particles”

Since 07/2010 Postdoc Institute Soft Matter and Functional Materials, EU-Project POLYCAT (representative of the workpackage leader), maintaining industry contacts, Helmholtz-Zentrum Berlin, Germany, Prof. Dr. M. Ballauff

Scholarships

10/2002 – 08/2007 Hochbegabtenstipendium nach BayBFG

05/2008 – 04/2010 Graduiertenstipendium nach BayEFG

Selected Publications

- Hoffmann, M. *Synthesis and Characterization of Anisotropic Colloidal Particles*; Logos Berlin: Berlin, 2010.
- Hoffmann, M.; Siebenbürger, M.; Harnau, L.; Hund, M.; Hanske, C.; Lu, Y.; Wagner, S.; Drechsler, M.; Ballauff, M.: Thermoresponsive Colloidal Molecules. *Soft Matter*, **2010**, 6, 1125.
- Hoffmann, M.; Wagner, C. S.; Harnau, L.; Wittemann, A.: 3D Brownian Diffusion of Submicron-Sized Particle Clusters. *ACS NANO*, **2009**, 3, 3326.
- Hoffmann, M.; Jusufi, A.; Schneider, C.; Ballauff, M.: Surface potential of spherical polyelectrolyte brushes in the presence of trivalent counterions. *J. Coll. Interf. Sci.* **2009**, 338, 566.
- Hoffmann, M.; Lu, Y.; Schrinner, M.; Ballauff, M.; Harnau, L.: Dumbbell-Shaped Polyelectrolyte Brushes Studied by Depolarized Dynamic Light Scattering. *J. Phys. Chem. B.* **2008**, 112, 14843.

Dr. Yan Lu

(1976)

Nationality: P. R. China

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Tel.: +49 30 8062 43191; Fax: +49 30 8062 42308
Email: yan.lu@helmholtz-berlin.de



Research interest

- Polymer colloids, including polyelectrolyte brushes and microgels
- Organic/inorganic hybrid colloid particles and their application as catalyst, sensor and solar cells

University Education

09.1994 – 07.1998 Bachelor in Polymer Science and Engineering in College of Material Science and Engineering, Donghua University, Shanghai, P. R. China

09.1998 – 04.2001 M.S. in Material Science in College of Material Science and Engineering, Donghua University, Shanghai, P. R. China

Scientific Degrees

Dr. rer. nat.: Chemistry, in Institute for Macromolecular Chemistry and Textile Chemistry, Dresden University of Technology, Germany
(02. 2005)
Supervisor: Prof. Dr. H. J. P. Adler

Scientific Development

04.2005 – 08.2006 Postdoc in the group of Prof. Dr. M. Ballauff, Physical Chemistry I, University of Bayreuth, Germany

09.2006 - 08.2009 Akademische Rätin, Physical Chemistry I, University of Bayreuth, Germany

Since 09.2009 Group leader of Colloid Chemistry, Institute of Soft Matter and Functional Materials, Helmholtz-Zentrum Berlin für Materialien und Energie, Germany

Selected Publications

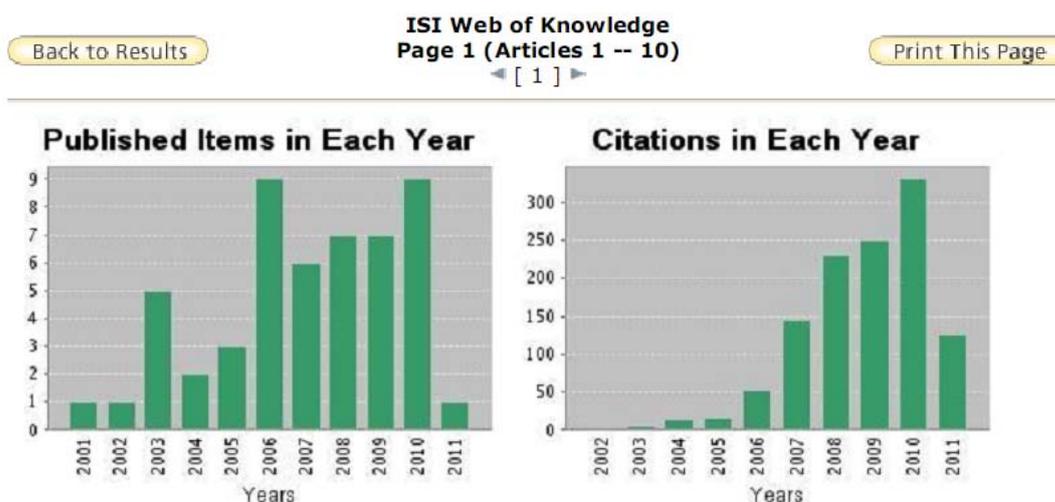
1. Y. Lu, Y. Mei, M. Ballauff*, M. Drechsler, Thermosensitive Core-Shell Particles as Carriers for Ag Nanoparticles: Modulating the Catalytic Activity by a Phase Transition in Networks, *Angewandte Chemie Int. Ed.* **45**, 813 (2006).
2. M. Ballauff, Y. Lu*, "Smart" Nanoparticles: Preparation, Characterization and Applications, *Polymer (Feature Article)* **48**, 1815 (2007).
3. Y. Lu*, S. Proch, M. Schrunner, M. Drechsler, R. Kempe, M. Ballauff, Thermosensitive Core-Shell Microgel as a "Nanoreactor" for Catalytic Active Metal Nanoparticles, *J. Mater. Chem.* **19**, 3955 (2009).
4. R. Sai Yelamanchili, Y. Lu*, T. Lunkenbein, N. Miyajima, L. Yan, M. Ballauff, J. Brey, Shaping colloidal rutile into thermally stable and porous mesoscopic titania-balls, *Small* **5**, 1326 (2009).
5. Y. Lu*, J. Yuan, F. Polzer, M. Drechsler, J. Preussner, In-situ Growth of Catalytic Active Au-Pt Bimetallic Nanorods in Thermo-Responsive Core-Shell Microgels, *ACS Nano* **4**, 7078 (2010).

Patents

A. Karpov, H. Hibst, A. Terrenoire, A. Weiss, M. Ballauff, Y. Mei, Y. Lu, R. Kempe, S. Proch, J. Villanueva, Technique of cross-coupling reactions with metallic nanoparticles as catalyst, WO 2008/074702 A1.

Award

2005: "APi-Prize" as the best dissertation in 2005 by the German Chemical Society (GDCh) Division of Coatings and Pigments.



Results found: 51

Sum of the Times Cited: 1,168

Average Citations per Item: 22.90

h-index: 19

Dr. Uwe Müller

(1967)

Macromolecular crystallography group at BESSY-II
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Area of Expertise and Field of Interest

Area of Expertise: Macromolecular crystallography using synchrotron radiation
Field of Interest: Protein structure and functions, Instrumentation of X-ray diffraction
beamlines

Scientific Development

Study 1989-1994 Study of Chemistry at the Humboldt University in Berlin
PhD 1999 at Free University Berlin (Prof. Dr. U. Heinemann)
Post-doc 1999-2003 Institut für Kristallographie, Free University Berlin, Prof.
Dr. W. Saenger
2004- 2008 Staff scientist and group leader, BESSY-GmbH, Berlin
Since 2009 Staff scientist and group leader, Helmholtz Zentrum Berlin

Invited Talks (selection)

"3th Winter school for soft X-ray in macromolecular crystallography", Berlin 2009
"Synchrotron radiation instrumentation 2009", Melbourne 2009

Selected publications

- [1] Perl, D., U. Mueller, et al. (2000). *"Two exposed amino acid residues confer thermostability on a cold shock protein."* Nat Struct Biol 7(5): 380-3.
- [2] Mueller, U., L. Nyarsik, et al. (2001). *"Development of a technology for automation and miniaturization of protein crystallization."* J Biotechnol 85(1): 7-14.
- [3] Heinemann, U., K. Bussow, Mueller U., et al. (2003). *"Facilities and methods for the high-throughput crystal structural analysis of human proteins."* Acc Chem Res 36(3): 157-63.
- [4] Schonfeld, D. L., R. B. Ravelli, Mueller U., et al. (2008). *"The 1.8-Å crystal structure of alpha1-acid glycoprotein (Orosomuroid) solved by UV RIP reveals the broad drug-binding activity of this human plasma lipocalin."* J Mol Biol 384(2): 393-405.
- [5] Klein, N., I. Senkovska, Mueller U., et al. (2009). *"A mesoporous metal-organic framework."* Angew Chem Int Ed Engl 48(52): 9954-7.

Priv. Doz. Dr. Gerd Schneider

(1963)

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Area of Expertise and Field of Interest

Area of Expertise: X-ray physics, X-ray imaging, Fourier and X-ray optics,
nanotechnology, synchrotron radiation and biophysics

Field of Interest: X-ray imaging, nano-tomography and X-ray spectromicroscopy
applied to biological cells and IT-devices

Scientific Development

Study	1983 - 1988 Study of chemistry at the University of Göttingen
PhD	1992 University of Göttingen (Prof. Dr. G. Schmahl)
	1992 Ernst-Eckhard-Koch award
Post-doc	1992 - 1995 Institute for X-ray Physics, University of Göttingen,
	1996 - 1999 Research Assistant (C1), Institute for X-ray Physics, University of Göttingen
	1999 Habilitation in Physics, University of Göttingen
	2000 Heisenberg Fellowship of the DFG
Senior Scientist	2000 - 2002 Lawrence Berkeley National Laboratory
	2003 - 2008 Group leader, BESSY m.b.H., Berlin
	2005 Priv.Do. at Humboldt Universität zu Berlin
	2009 Head of Department for Microscopy, Helmholtz Zentrum Berlin
	2010 Group leader Microscopy Helmholtz Zentrum Berlin

Selected Scientific Expert, Review or Council Activities, Awards

Ernst-Eckhard-Koch award 1992, Heisenberg fellowship 2000, Editorial Advisory Board of Current Nanoscience (since 2005), Member of the International Program Committee of the International Conference on X-ray Microscopy (since 2005), Reviewer for Nature, Applied Physics Letters, Ultramicroscopy, Optics Express, referee for the Körber-Stiftung

Invited Talks (selection)

2009 NSLS/CFN Joint Users' Meeting, Workshop 2: The Cold, Soft Truth: Cryo Systems for Studying Soft Materials, Brookhaven National Laboratory, May 18, 2009

SCANDEM 2009-Annual Meeting of the Nordic Microscopy Society University of Iceland, Reykjavík, Iceland, 2009

First International Symposium on Structural Systems Biology, Hamburg, September 24- 25, 2009

Frontiers in Optics, San Jose, USA 2009

Seminar at SOLEIL Synchrotron, November 3, 2009, France

Workshop on Correlative Microscopy, Oxford 2010; England

Microscience 2010, London 2010;

“3rd International Workshop on Imaging Techniques with Synchrotron Radiation, Suzhou 2010, China

Selected publications

- [1] Ehrenfried Zschech, Rene Huebner, Dmytro Chumakov, Oliver Auel, Daniel Friedrich, Peter Guttman, Stefan Heim, Gerd Schneider, *Stress-induced phenomena in nanosized copper interconnect structures studied by x-ray and electron microscopy*, J. Appl. Phys. **106**, 093711 (2009)
- [2] José L. Carrascosa, Francisco Javier Chichón, Eva Pereiro, María Josefa Rodríguez, José Jesús Fernández, Mariano Esteban, Stefan Heim, Peter Guttman, Gerd Schneider, *Cryo-x-ray tomography of Vaccinia Virus membranes and inner compartments*, J. Struct. Biol. **168** (2009), 234-239; 186 (2009) 012041
- [3] G. Schneider, S. Rehbein, S. Werner, *Volume Effects in Zone Plates* in: Modern Developments in X-Ray and Neutron Optics Springer Series in Optical Sciences, Springer Berlin/Heidelberg **137** (2008), 137-171
- [4] S. Rehbein, S. Heim, P. Guttman, S. Werner, G. Schneider, *Ultra-high-resolution soft-x-ray microscopy with zone plates in high orders of diffraction*, Phys. Rev. Lett. **103**, (2009) 110801
- [5] G. Schneider, P. Guttman, S. Heim, S. Rehbein, F. Mueller, K. Nagashima, J.B. Heymann, W.G. Müller, J.G. McNally, *Three-dimensional cellular ultrastructure resolved by X-ray microscopy*, Nature Methods **7** (2010), 985-987

Dr. Sebastian Seiffert (Dipl.-Chem.)

(1979)

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Email: sebastian.seiffert@helmholtz-berlin.de
seiffert@chemie.fu-berlin.de



Area of Expertise

Polymer Chemistry, Polymer Physics, Physical Chemistry, Chemical Engineering

Scientific Career

- Since 01/2011 Leader of an independent research group at Helmholtz-Zentrum Berlin and Free University Berlin, Germany
- Funded by a Liebig Grant; Fund of the German Chemical Industry
 - Research on *Supramolecular Polymer Gels as Functional Materials*
 - Lecturer on *Polymer Physics* at Free University Berlin
- 01/2009–12/2010 Postdoctoral fellow in the group of Prof. D. A. Weitz, Dpt. of Physics and SEAS, Harvard University, Cambridge, Massachusetts, U.S.A.
- Research Fellowship by the German Acad. of Sciences Leopoldina
 - Subject of Research: *Functional Polymer Microgels*
- 01/2008–12/2008 Postdoctoral fellow in the group of Prof. W. Oppermann, Institute of Physical Chemistry, Clausthal University of Technology, Germany
- Subject of Research: *Structure and Dynamics in Polymer Gels*
 - Lecturer on “*Structure and Dynamics in Polymer Systems*”
- 12/2007 PhD degree, Clausthal University of Technology, Germany
- Predicate: summa cum laude
 - Thesis “*Structure and Tracer Dynamics in Polyacrylamide Gels*”
- 08/2004–12/2007 PhD student in the group of Prof. W. Oppermann, Institute of Physical Chemistry, Clausthal University of Technology, Germany
- Subject of research: *Structure and Dynamics in Polymer Hydrogels*
- 07/2004 Diploma in Chemistry, Clausthal University of Technology
- Degree: Dipl.-Chem; passed with distinction (mit Auszeichnung)
 - Diploma thesis “*Diffusion of Linear Polyacrylamide Chains in Semidilute Systems*”
- 10/1999–08/2004 Study of Chemistry at Clausthal University of Technology

Awards and Grants

- Since 01/2011 "Liebig Scholarship" by the Fund of the German Chemical Industry for the establishment of an independent junior research group at the Helmholtz Center for Materials and Energy Berlin and FU Berlin
- 01/2009–12/2010 Scholarship by the German Academy of Sciences Leopoldina for a biennial postdoctoral stay at the Department of Physics and SEAS, Harvard University, Cambridge, Massachusetts, U.S.A.
- 10/2009 Dissertation Award by the Society of Friends of Clausthal University of Technology
- 06/2008 Offer of a Postdoc Scholarship by the German Research Foundation (declined in favor of scholarship by the German Academy of Sciences Leopoldina)
- 02/2002 Nominee for scholarship by the German National Academic Foundation
- 12/2001 Book award of the local section "Harz" of the German Chemical Society (GDCh) for outstanding performances as an undergraduate

Selected Publications

- S. Seiffert and D. A. Weitz, "Controlled Fabrication of Polymer Microgels by Polymer-Analogous Gelation in Droplet Microfluidics." *Soft Matter* **2010**, *6*, 3184–3190.
- S. Seiffert, J. Thiele, A. R. Abate and D. A. Weitz, "Smart Microgel Capsules from Macromolecular Precursors." *J. Am. Chem. Soc.* **2010**, *132*, 6606–6609.
- S. Seiffert and W. Oppermann, "Diffusion of Linear Macromolecules and Spherical Particles in Semidilute Polymer Solutions and Polymer Networks." *Polymer* **2008**, *49*, 4115–4126.
- S. Seiffert, W. Oppermann and K. Saalwächter, "Hydrogel Formation by Photocrosslinking of Dimethylmaleimide Functionalized Polyacrylamide." *Polymer* **2007**, *48*, 5599–5611.
- S. Seiffert and W. Oppermann, "Systematic Evaluation of FRAP Experiments Performed in a Confocal Laser Scanning Microscope." *J. Microsc.* **2005**, *220*, 20–30.

Dr. Roland Steitz

(1960)

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Energie GmbH, Hahn-Meitner-Platz 1, 14109 Berlin

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Area of Expertise and Field of Interest

Area of Expertise: Physics of soft matter interfaces

Field of Interest: Chemistry and physics of amphiphiles, scattering methods

Scientific Development

Study 1978- 1988 Study of chemistry at the University of Mainz

PhD 1993 at the University of Mainz (Prof. Dr. H. Möhwald)

Post-doc 1994-1995 Dept. Chemistry, University of Queensland, Brisbane,
Australia, Dr. G. Barnes

1995- 2000 Junior Scientist, Technische Universität Berlin

2000- 2004 Junior Scientist, Max-Planck Institut für Kolloid und
Grenzflächenforschung, Golm

Senior Scientist Since 2004, Institute for Soft Matter and Functional Materials, HZB,
Berlin

Habilitation at TU Berlin in progress

Selected Scientific Expert, Review or Council Activities, Awards

Member of Scientific Panel: Sub-Committee 9, Institut Laue-Langevin, Grenoble, France
(2003-2006), Member of BENSC Scientific Panel, Hahn-Meitner-Institut, Berlin (2003),
Member of German Committee for Research with Neutrons, KFN (since 2005)

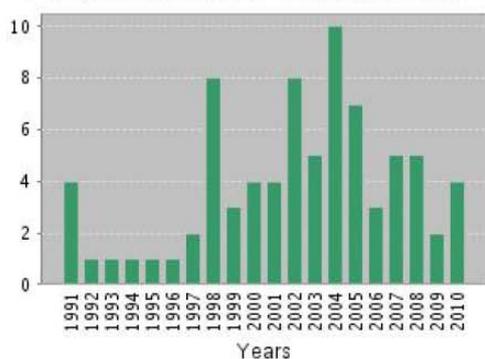
Invited Talks (selection)

“Physikalisches Kolloquium”, Universität des Saarlandes, Saarbrücken 2005; “52nd AVS
International Symposium”, Boston, 2005; “ADAM Workshop” Ruhr-Universität, Bochum 2006;
“2nd BENSC Adsorption Workshop HMI, Berlin 2007; „Workshop zum Thema Streumethoden,
SPP 1273 Kolloidverfahrenstechnik“, Bayreuth 2008; “GISAS 2009”, Satellite Conference of
SAS 2009, Hamburg 2009; Department of Physics and Astronomy, Uppsala University,
Sweden 2010; “DPG-Frühjahrstagung”, TU-Dresden, 2011

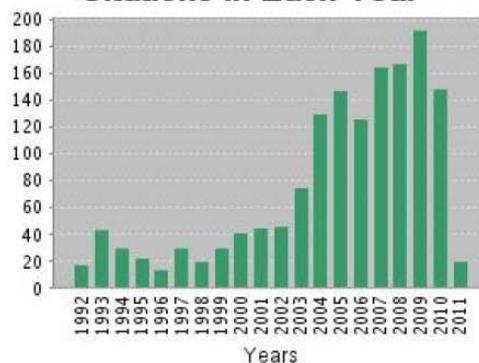
Selected publications

1. M. Kreuzer, T. Kaltofen, R. Steitz, B. H. Zehnder, R. Dahint, *Pressure cell for investigations of solid-liquid interfaces by neutron reflectivity*, Review of Scientific Instruments 2011, **82** (2), 023902-7
2. F. Evers, R. Steitz, M. Tolan, C. Czeslik, *Analysis of Hofmeister effects on the density profile of protein adsorbates - a neutron reflectivity study*, Journal of Physical Chemistry B 2009, **113**, 8462-8465
3. M. Wolff, R. Steitz, P. Gutfreund, N. Voss, S. Gerth, M. Walz, A. Magerl, H. Zabel, *Shear Induced Relaxation of Polymer Micelles at the Solid-Liquid Interface*, Langmuir 2008, **24**, 11331
4. V. Papaefthimiou, R. Steitz, G. H. Findenegg, *Schaltbare Oberfläche – Responsive Polymerschichten* Chemie in unserer Zeit 2008, **42**, 102-115
5. C. Czeslik, G. Jackler, R. Steitz and H.-H. von Grünberg, *Protein Binding to Like-Charged Polyelectrolyte Brushes by Counterion Evaporation*, J. Phys. Chem. B 2004, **108**, 13395.
6. R. Steitz, T. Gutberlet, T. Hauß, B. Klösgen, R. Krastev, S. Schemmel, A. C. Simonsen and G. H. Findenegg, *Nanobubbles and Their Precursor Layer at the Interface of Water Against a Hydrophobic Substrate*, Langmuir 2003, **19**, 2409
7. R. Steitz, W. Jaeger, R. v. Klitzing, *Influence of charge density and ionic strength on the multilayer formation of strong polyelectrolytes*, Langmuir 2001, **17**, 4471
8. R. Steitz, V. Leiner, R. Siebrecht and R. v. Klitzing, *Influence of the ionic strength on the structure of polyelectrolyte films at the solid/liquid interface*, Colloids and Surfaces A 2000, **163**, 63
9. M. Tarabia, H. Hong, D. Davidov, S. Kirstein, R. Steitz, R. Neumann, Y. Avny, *Neutron and X-ray reflectivity studies of self-assembled heterostructures based on conjugated polymers*, J. Appl. Phys. 1998, **83**, 725-732

Published Items in Each Year



Citations in Each Year



Results found: 83

Sum of the Times Cited: 1,534

Average Citations per Item: 18.48

h-index: 22

Dr. Manfred S. Weiss (1963)

Macromolecular crystallography group at BESSY-II
F-I2 Soft Matter and Functional Materials
Helmholtz-Zentrum Berlin für Materialien und Energie GmbH
Albert-Einstein-Str. 15
D-12489 Berlin

Tel: 030 8062-13149 (-14975, Fax)
Email: manfred.weiss@helmholtz-berlin.de



Area of Expertise and Field of Interest

Area of Expertise: Macromolecular crystallography using synchrotron radiation
Field(s) of Interest: Protein structure and function, macromolecular crystallization,
instrumentation of X-ray diffraction beamlines

Scientific Development

Study 1982-1989: Study of Chemistry at the Albert-Ludwigs-Universität Freiburg
i. Br. and the University of Massachusetts at Amherst, USA.
Ph.D. 1989-1992: at Albert-Ludwigs-Universität Freiburg (Prof. Dr. G. E. Schulz)
Post-doc 1992-1996 Molecular Biology Institute, University of California at Los Angeles,
USA (Prof. Dr. D. Eisenberg)
1996-2001 Senior Research Assistant at the Institute for Molecular
Biotechnology, Jena (Prof. Dr. R. Hilgenfeld)
2001-2009 Team Leader at the EMBL Hamburg Outstation
2009-present Staff scientist, Helmholtz Zentrum Berlin

Invited Talks (selection)

Conference: European Crystallographic Meeting ECM-26, 31.08.-02.09.2010, Darmstadt,
Germany
Conference: American Crystallographic Association (ACA) Meeting 2010, 24.-30.07.2010,
Chicago, USA
Course: *Training Methods for Macromolecular Crystallography. From Measurement to Model -
2009, M2M-9, 21.-28.10.2009, EMBL Hamburg Outstation, Hamburg,
Germany.*
EMBO-Course: *Exploiting Anomalous Scattering in Macromolecular Structure Determination,*
15.-22.06.2009, ESRF Grenoble, France.
3rd Winter School on *Soft X-rays in Macromolecular Crystallography*, 18.-20.02.2009, Berlin,
Germany.
Conference: International Union of Crystallography (IUCr) Conference 2008, 23.-30.08.2008,
Osaka, Japan.
Workshop: ISRTMSF2008, 07.-11.01.2008, University of Madras, Guindy Campus, Chennai,
India.

Selected publications

- T. Werther, A. Zimmer, G. Wille, R. Golbik, **M. S. Weiss** & S. König (2010). New Insights into Structure-Function Relationships of Oxalyl-CoA Decarboxylase from *Escherichia coli*. *FEBS J.* 277, 2628-2640.
- M. J. Belousoff, C. Davidovich, E. Zimmerman, Y. Caspi, I. Wekselman, L. Rozenszajn, T. Shapira, O. Sade-Falk, L. Taha, A. Bashan, **M. S. Weiss** & A. Yonath (2010). Ancient Machinery Embedded in the Contemporary Ribosome. *Biochem Soc Trans.* 38, 422-427.
- R. Janowski, G. Kefala & **M. S. Weiss** (2010). The structure of Dihydrodipicolinate Reductase (DapB) from *Mycobacterium tuberculosis* in Three Crystal Forms. *Acta Cryst.* D66, 61-72.
- L. Schuldt, S. Weyand, G. Kefala & **M. S. Weiss** (2009). The Three-dimensional Structure of a Mycobacterial DapD Provides Insights into DapD Diversity and Reveals Unexpected Particulars About the Enzymatic Mechanism. *J. Mol. Biol.* 389, 863-879.
- R. Janowski, S. Panjikar, A. Nasser Eddine, S. H. E. Kaufmann & **M. S. Weiss** (2009). Structural Analysis Reveals DNA binding Properties of Rv2827c, a Hypothetical Protein from *Mycobacterium tuberculosis*. *J. Struct. Funct. Genom.* 10, 137-150.
- C. Mueller-Dieckmann, S. Panjikar, A. Schmidt, S. Mueller, J. Kuper, A. Geerlof, M. Wilmanns, R. K. Singh, P. A. Tucker & **M. S. Weiss** (2007). On the Routine Use of Soft X-Rays in Macromolecular Crystallography, Part IV - Efficient Determination of Anomalous Substructures in Bio-Macromolecules Using Longer X-ray Wavelengths. *Acta Cryst.* D63, 366-380.
- C. Mueller-Dieckmann, S. Kernstock, M. Lisurek, J. P. von Kries, F. Haag, **M. S. Weiss** & F. Koch-Nolte (2006). The Structure of Human ADP-Ribosylhydrolase 3 (ARH3) Provides Insights into the Reversibility of Protein ADP-Ribosylation. *Proc. Natl. Acad. Sci. USA* 103, 15026-15031.

Soft Matter and Functional Materials

1. **Research on Soft Matter and Functional Materials at the Helmholtz-Zentrum Berlin für Materialien und Energie**.....

The Institute of Soft Matter and Functional Materials

Organization of the Institute

Beamlines and Laboratories of the Institute

Soft Matter and Functional Materials: Research and Recent Highlights

Statistics User Service

Organizational Chart

2. **Beamlines**.....

V1

V6

V16

V18

BL 14.1

BL 14.2

BL 14.3

U41 X-ray microscope

Electron beam writer

3. **Laboratories**.....

BioLab

Chemistry Lab

Colloid Physics Lab

Joint Laboratory for Structural Research (IRIS)

Joint MX lab

Laboratory for Microfluidics

4. **Research**.....

Macromolecular Crystallography (MX)

Biophysics

Colloid Physics

Interfaces

Colloid Chemistry

Soft Matter Theory

X-Ray Microscopy

Polymer Physics

5. **CVs**.....

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