## Annette H. Erbse, Ph.D.

## Director of the Biochemistry Shared Instruments Pool • Director of the Macromolecular X-Ray Core

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| Professional Career: |   |
|----------------------|---|
| 2016-present         | Director of the Macromolecular X-ray core in the Department of Chemistry and Biochemistry, JSC Biotechnology Building, University of Colorado at Boulder  |
|                      | Responsibilities: Training and supervision of users; radiation license, trouble shooting of experiments and instruments, instrument maintenance; organisation and support of data collection at synchrotrons, administration and accounting for the core  |
| 2012-present         | Director of the Shared Instrument Core in the Department of Chemistry and Biochemistry, JSC Biotechnology Building, University of Colorado at Boulder   |
|                      | Responsibilities: Training and supervision of users; support in project development, experimental design, optimisation and evaluation; trouble shooting of experiments and instruments, instrument maintenance; lead in shared instrument grant proposals; service measurements for in house and commercial users; administration and accounting for the core |
| 2009-2012            | <b>Senior Research Associate,</b> University of Colorado, Boulder in the group of Prof. J.J.Falke., Boulder, CO, USA  |
|                      | Research project: Investigating the mechanisms of bacterial chemotactic signal sensing  |
| 2009-present         | <b>Manager EPR facility</b> Department of Chemistry/Biochemistry, University of Colorado – Boulder, Boulder, CO, USA  |
| 2006-2009            | <b>Research Associate,</b> University of Colorado, Boulder in the group of Prof. J.J.Falke, Boulder, CO, USA  |
|                      | Research project: Investigating the mechanisms of bacterial chemotactic signal sensing  |
| 2002-2005            | <b>Oberassistent (Senior Staff Scientist)</b> at the Centre for Molecular Biology, University of Heidelberg in the group of Prof. B. Bukau, Heidelberg, Germany   |
|                      | Research project: Function and mechanism of molecular chaperons in proteolysis  |
| 1999-2002            | <b>Post-doctoral Fellow</b> of the Howard Hughes Medical Institute at Yale University in the group of Prof. A. L. Horwich, New Haven, CT, USA   |
|                      | Research Project: Investigating the structure of TTR amyloid-fibers using EPR-spectroscopy  |
| 1996 -1999           | <b>Research Associate</b> , Department of Biological Sciences, University of Birmingham in the group of Dr. P. A. Lund, Birmingham, England   |

Annette Erbse Curriculum Vitae Research project: Function and mechanism of molecular chaperons of the HSP60 family **Education:** October 1996 PhD at the Department of Chemistry, University of Kaiserslautern, Germany 1993 - 1996PhD student in the group of Prof. W. Trommer und Dr. P. Vogel at the Department of Chemistry, University of Kaiserslautern, Germany Thesis project: Investigating the structure – function relationship of different ATP binding proteins employing EPR-spectroscopy February 1993 Diploma in Chemistry at the University of Kaiserslautern, Germany (equivalent to Masters in the USA) 1992 - 1993**Diploma Thesis** (equivalent to **Masters Thesis**) at the University of Kaiserslautern, Department of Chemistry in the group of Prof. W. Trommer Thesis project: Kinetic investigations on native and activated F<sub>1</sub>-ATPase from chloroplasts

## **Expertise:**

1986 - 1992

- Developing biophysical/biochemical assays for the analyses of protein, DNA and RNA structure and function.
- Protein, DNA and RNA characterization. SDS gel-electrophoresis, western blotting, activity assays, mass spectrometry, Fluorescence- and EPR spectroscopy, calorimetric assays, circular dichroism, dynamic and static multi angle light scattering, microscale thermophoresis.

Chemistry student at the Department of Chemistry at the University of Kaiserslautern,

- Enzyme kinetics and thermodynamics. Coupled activity assays, radioactive labelled substrates, different spectroscopic methods (EPR, fluorescence, FRET, anisotropy, UV-VIS) in steady-state or stopped-flow experiments), SPR (Biacore), HD exchange, peptide libraries, fast kinetics (stopped-flow and chemical quench flow), Isothermal Titration Calorimetry, Microscale Thermophoresis, investigation of cooperativity and analyses of data by applying mathematical models.
- Macromolecular X-ray Crystalography

Germany

- Protein mass spectrometry. MALDI and ESI-LC/MS, MS/MS, accurate total mass determination, peptide fingerprinting, PDS, Instruments: ABI Q-Star Pulsar, Bruker Ultraflex MALDI-TOF/TOF and Voyager-DE™ STR). Mass spectrometry teaching: Co-developed and team-taught a graduate mass spectrometry hands-on/theory class with Dr. M.M. Mayer (Center for Molecular Biology Heidelberg).
- Expression and purification of recombinant proteins. Purification employing HPLC und FPLC methods (gel filtration, ion exchange -, reversed-phase- and affinity chromatography), purification from inclusion bodies, purification of radioactively labelled proteins.
- Developing structure/function guided protein mutations.
- Protein modification. Site-specific biotinylation, PEGylation, spin labelling or fluorophore conjugation employing designed, structure/function guided cysteine mutants via maleimide, succinimide, disulfide chemistries, genetically encoded affinity tags, enzymatic modification and chemical modification.
- Isolation of membrane fractions, inner membrane vesicles, trans-membrane receptors and membranebound, multi-protein complexes in native membranes.
- Molecular biology (sub-cloning, mutagenesis, constructing fusion proteins).