Little Grassy Lake Conference Center Southern Illinois University Carbondale, Illinois

Organized by

Kenneth P. Murphy University of Iowa College of Medicine

and

Michael D. Brenowitz Albert Einstein College of Medicine

October 7 -10, 1995





 ◆ 4:00 pm Registration at Main Desk, Little Grassy Lodge

• 7:00 pm Wine and Cheese Reception

No dinner will be served.

Posters for Session I (A - La by last name of the first author) may be set up in the *Sledgefoot Lodge* (adjacent to the dining room, *Freeberg Hall*) Saturday night and Sunday morning.

Posters for Session II (Li - Z by last name of the first author) may be set up in the *Sledgefoot Lodge* (adjacent to the dining room, *Freeberg Hall*) Monday morning.



Sunday October 8

♦ 7:00 am Breakfast, Freeberg Hall

♦ 8:30 am Keynote Address I

Victor Bloomfield University of Minnesota

DNA Condensation and Intermolecular Forces

Session I: Electrostatics

♦ 9:30 am Refreshments

♦ 9:50 am Kim Sharp

University of Pennsylvania

Determination of the Heat Capacities of Polar and Apolar Aqueous Solutions Using the Random Network Model (RNM)

♦ 10:35 am Bertrand Garcia-Moreno

The Johns Hopkins University

Experimental Studies of Electrostatic Interactions in Proteins

◆ 11:20 am Katrina Wagner Vanderbilt University

Structure in the Electric Potential of DNA

• 11:45 am M. Thomas Record, Jr.

University of Wisconsin, Madison

lon Effects on Interactions of Two Charged Ligands KWK₈NH₂ (+8) and Εσ⁷⁰RNA

Polymerase) with DNA

◆ 12:30 pm Lunch, Freeberg Hall

Session II: Nucleic Acids

• 3:00 pm David Draper

The Johns Hopkins University

Protein, Antibiotic and Ion Binding to a Conserved Ribosomal RNA Fragment

• 3:45 pm Girija Krishnamurthy

Wyeth Ayerst Research

Recent Biophysical and Biological Studies on the Mechanism of Action of Calicheamicin

♦ 4:10 pm Refreshments

♦ 4:30 pm Vitaly Buckin

New York University

Hydration Effects Upon Substituting Na⁺ for Cs⁺ or Mg²⁺ in the Ionic Atmosphere of poly (rA)•2poly (rU) and poly (rA)•poly (rU) Helicles

♦ 4:55 pm Keith Bjornson

Washington University School of Medicine

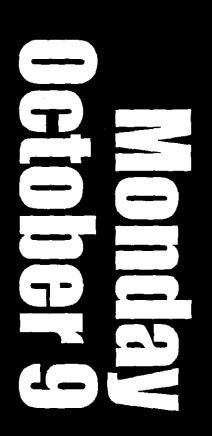
Kinetic Mechanism of DNA Binding and DNA-Induced Dimerization of the *Escherichia coli* Rep Helicase

♦ 5:20 pm Kathleen Hall

Washington University School of Medicine

Thermodynamics of the Association of the UIA RNA Binding Domain and RNA Hairpins

♦ 6:35 pm Dinner, Buffalo Tro at Freeberg Hall



♦ 7:00 am

Breakfast, Freeberg Hall

♦ 8:30 am

Keynote Address II

Mario Amzel

The Johns Hopkins University

Binding of Flexible Peptides: Structure and Thermodynamics. Calculation of Entropy

Session III: Linkage & Allostery

♦ 9:30 am

Refreshments

♦ 9:50 am

Donald Senear

The University of California, Irvine

Coordinate Gene Regulation in E. coli: Competitive and Inducer Controlled

Cooperative Interactions between CAP and

CytR at the deoP2 Promoter.

♦ 10:35 am

Michael Dovle

Smith-Kline Beechem Pharmaceuticals

Inhibition Mechanism of Beta Trypsin by

Ecotin

◆ 11:20 am Yingwen Huang

Washington University School of Medicine

Transformation of Cooperative Free Energies

(⁸∆G_c) between Ligation Systems of Hemoglobin: An Exact Resolution of the **Carbon Monoxide Binding System**

• 11:45 am J. Ching Lee

Univ. Texas Medical Center, GalvEston

The Modus Operandi of E.Coli CRP

12:30 pm Lunch, Freeberg Hall

Session IV: Proteins

♦ 3:00 pm

Ernesto Freire

The Johns Hopkins University

Thermodynamics of Protein Stability and

Molecular Recognition

♦ 3:45 pm

Sydney D. Hoeltzli

Washington Univ. School of Medicine

Stopped-Flow Spectroscopy: Real-time Unfolding Studies of 6-19F-Tryptophan labeled

E. coli Dihydrofolate Reducatase

♦ 4:10 pm

Refreshments

♦ 4:30 pm

Nadine C. Gassner

University of Oregon

Generic Core Replacement in T4 Lysozyme

Preserves Protein Activity, Folding

Cooperativity & Shape

♦ 5:15 pm

Peter L. Privalov

The Johns Hopkins University

On the Entropy of Protein Folding

♦ 6:30 pm

Dinner, Freeberg Hall



♦ 7:00 am Breakfast, Freeberg Hall

Session V: Molecular Recognition

+ 8:20 am Ruth Saecker Spolar

University of Wisconsin, Madison

Thermodynamic Signatures of Conformational Changes in Protein-Nucleic Acid Interactions

♦ 9:05 am Javier Gómez

The Johns Hopkins University

Thermodynamic Mapping of the Inhibitor Site of the Aspartic Proteinase Endothiapepsin

♦ 9:30 am Refreshments

♦ 9:50 am Patrick S. Stayton

University of Washington

Molecular Origins of High-Affinity and Slow Dissociation Kinetics in Protein-Ligand Interactions

♦ 10:15 am Patrick Connelly

Groton School

Immunosuppressive Drug Design and Inosine Monophosphate Dehydrogenase: A Proposed Solution to a Problem in *Differential Molecular Recognition*

♦ 11:00 am Lunch, Freeberg Hall



Lectures

DNA Condensation and Intermolecular Forces

Victor Bloomfield

Determination of the Heat Capacities of Polar and Apolar Aqueous Solutions Using the Random Network Model (RNM)

Kim Sharp & Bhupinder Madan

Experimental Characterization of Electrostatic Forces in Proteins

Bertrand García-Moreno E.

Structure in the Electric Potential of DNA

Katrina Wagner, Thomas Kephart, David Hochberg and Glenn Edwards

Ion Effects on Interactions of Two Charged Ligands (KWK6NH2(+8) and E σ 70 RNA Polymerase) with DNA

M. Thomas Record, Jr.

Protein, Antibiotic and Ion Binding to a Conserved Ribosomal RNA Fragment D.E. Draper, S. Huang & Y. Xing

Recent Biophysical and Biological Studies on the Mechanism of Action of Calicheamicin

Girija Krishnamurthy and George A. Ellestad

Hydration Effects Upon Substituting Na⁺ for Cs⁺ or Mg²⁺ in the Ionic Atmosphere of poly (rA)•2poly (rU) and poly (rA)•poly (rU) Helicies

Vitaly Buckin & Luis A. Marky

Kinetic Mechanism of DNA Binding and DNA-Induced Dimerization of the Escherichia coli Rep Helicase

Keith P. Bjornson, Keith J.M. Moore and Timothy M. Lohman

Thermodynamics of the Association of the UIA RNA Binding Domain and RNA Hairpins

Kathleen B. Hall

Binding of Flexible Peptides: Stucture and Thermodynamics. Calculation of Entropy Changes

K.H. Lee, Y. Pan, A. D'Aquino, K. Murphy, D. Xie, E. Freire and M. Amzel

Coordinate Gene Regulation in E. coli: Competitive and Inducer Controlled Cooperative Interactions between CAP and CytR at the deoP2 Promoter

L.T. Perini, L.A. Doherty, E. Werner and D.F. Senear

Binding Affinity of B-Trypsin and Ecotin Measured by Differential Scanning Calorimetry

Michael L. Doyle, Cherice Lee, Mary McGrath, and Robert Fletterick

Transformation of Cooperative Free Energies(^{ij} \(\triangle G_c \)) between Ligation Systems of Hemoglobin: An Exact Resolution of the Carbon Monoxide Binding System Yingwen Huang and Gary K. Ackers

The Modus Operandi of E. coli CRP

J. Ching Lee

Thermodynamics of Protein Stability and Molecular Recognition Ernesto Freire

Stopped-Flow NMR Spectroscopy: Real-time Unfolding Studies of 6-19F-Tryptophan labeled E. coli Dihydrofolate Reductase Sydney D. Hoeltzli and Carl Frieden

Generic Core Replacement in T4 Lysozyme Preserves Protein Activity, Folding Cooperativity and Shape

Nadine C. Gassner, Walter A. Baase and Brian W. Matthews

On the Entropy of Protein Folding

Peter L. Privalov

Thermodynamic Signatures of Conformational Changes in Protein-Nucleic Acid Interactions

R. Saecker Spolar and M. Thomas Record, Jr.

Thermodynamic Mapping of the Inhibitor Site of the Aspartic Proteinase Endothiapepsin

Javier Gómez and Ernesto Freire

Molecular Origins of High-Affinity and Slow Dissociation Kinetics in Protein-Ligand Interactions

Patrick S. Stayton, Ashutosh Chilkoti, Lisa Klumb and Vano Chu

Immunosuppressive Drug Design and Inosine Monophosphate Dehydrogenase: A Proposed Solution to a Problem in Differential Molecular Recognition Patrick Connelly

Posters

E. coli Helicase II Unwinds DNA by an Active Mechanism

Janid A. Ali and Timothy M. Lohman

Folding transitions coupled to hirudin binding to thrombin

Youhna M. Ayala, Alessandro Vindigni, Murad Nayal, Ruth S. Spolar, M. Thomas Record, Jr., and Enrico Di Cera

Thermodynamics of Protein-Protein Interactions: Binding of Turkey ovomucoid third domain to porcine pancreatic elastase

Brian M. Baker and Kenneth P. Murphy

Structural Characterization of Multiple Partially Folded States of a BPTI Analogue Elisar Barbar, George Barany and Clare Woodward

Fluorescence Derivative of E. coli RNA Polymerase Sigma Subunit in the -10 Recognition Domain of the Protein

Sandhya Callaci and Tomasz Heyduk

A Hidden Contribution to van't Hoff Enthalpy Estimates from Small Heat Capacity Changes

Jonathan B. Chaires

Effects of Conserved Residues on the REgulation of Rabbit Muscle Pyruvate Kinase Xiaodong Cheng, Robert H.E. Friesen and J. Ching Lee

Mutagenesis studies of the slow and fast forms of thrombin

Quoc D. Dang, Enriqueta R. Guinto, Thalia Farazi, Alessandro Vindigni, Youhna M. Ayala, and Enrico DiCera

The Magnitude of the Backbone Conformational Entropy Change in Protein Folding Jose Alejandro D'Aquino, Javier Gómez, Vincent J. Hilser, Kon Ho Lee, L. Mario Amzel and Ernesto Freire

Comparing Stability in Structurally Homologous Proteins with Little Sequence Identity

G.T. DeKoster & A.D. Robertson

Equilibrium DNA Binding of the Sac7d Protein from the Hyperthermophile Sulfolobus acidocaldarius: Fluorescence and Circular Dichroism Studies Stephen P. Edmondson, James McAfee and John W. Shriver

Multi-Dimensional Thermodynamic Study of the Unfolding of Staphylococcal Nuclease A and Its V66W Mutant

Maurice R. Eftink

Microtuble Assembly Dynamics with Pressure and IR Vibrational Perturbations Daniel Engh, Bruce Johnson, Glenn Edwards and Robley Williams

Intramolecular Electrostatic Interactions Accelerate Hydrogen Exchange in Diketopiperazine Relative to 2-Piperidone

William R. Forsyth and Andrew D. Robertson

Stability And Calcium Binding of Recombinant Fragments of Troponin C R.S. Fredricksen and C. Swenson

Thermodynamic Identification of Binding Hot Spots in Protein/Protein Recognition Ernesto Freire, Javier Gómez, Vincent J. Hilser, Craig R. Johnson, and Dong Xie

Pressure-Induced Protein Denaturation of Staphylococcal Nuclease Kelly J. Frye and Catherine A. Royer

An RNA Conformational Switch

T.C. Gluick & D.E. Draper

The Heat Capacity of Proteins

Javier Gómez, Vincent J. Hilser, Dong Xie and Ernesto Freire

Heme-Protein Interaction and Hemoglobin Stability Under High Hydrostatic Pressure Probed By Fluorescence Spectroscopy

Zygmunt Gryczynski, Jacek Lubkowski, Sabrina Beretta and Enrico Bucci

Energetics of Hydrogen Bonding in Proteins: A Model Compound Study Susan M. Habermann and Kenneth P. Murphy

The Binding of an Fab to Dimeric Human Interleukin-5: A Solution Interaction Analysis Using the Time Derivative Method to Interpret Sedimentation Velocity Data Preston Hensley, Carol C. Silverman, Dean E. McNulty, Michael L. Doyle, David G. Myszka, Terence G. Porter and Walter F. Stafford III

Protein Footprinting: Mapping Domains of *E. coli* RNA Polymerase α Subunit Involved in Enzyme Assembly

Ewa Heyduk, Tomasz Heyduk, Konstantin Severinov, Hong Tang and Richard H. Ebright

The Contribution of van der Waals Interactions to the Enthalpy of Folding and Binding

Vincent J. Hilser, Javier Gómez and Ernesto Freire

Global Analysis Of The Urea And Acid-Induced Unfolding Of Stapyloccocal Nuclease

Roxana M. Ionescu and Maurice R. Eftink

Fatty Acids Abolish High-Affinity Binding of Glyburide to Human Serum Albumin: Implications for Type II Diabetes

Michael G. Jakoby IV, David Song, Douglas F. Covey and David P. Cistola

Biochemical and Biophysical Characterizations of a Temperature Sensitive Mutant of *Escherichia coli* trp Repressor, L75F TrpR

Lihua Jin, June Fukayama & Jannette Carey

Structural Stability of Small Oligomeric Proteins

Craig R. Johnson and Ernesto Freire

Visualizing DNA Electric Fields: An Easy-to-Implement Standard

Edward Keyes, Thomas Kephart and Glenn Edwards

Heterotropic effects of a strong allosteric effector (IHP) on the O2 and CO binding properties of the hemoglobin partially liganded species

Laurent Kiger, Claude Poyart and Michael C. Marden

Oxygen Binding Studies of Fell/Coll Hybrid Hemoglobins

Alexandra Klinger and Gary K. Ackers

The Role of Tyrosines in RBD Stability and RNA-binding

James K. Kranz and Kathleen B. Hall

Solvents interactions in hemoglobin: role of associated water on the enthalpy of a series of crosslinked hemoglobins

Herman Kwansa, Anna Razynska, Zygmunt Gryczynski, Enrico Bucci

mRNA Iron Responsive Element Thermodynamic Stability and NMR Structure Study Lance G. Laing and Kathleen B. Hall

K⁺-Ribosome Interactions Determine the Large Enhancements of 39K NMR Transverse Relaxation Rates in Viable Cells

M. Li, H.J. Guttman, S. Cayley, C.F. Anderson and M. Thomas Record, Jr.

CD and Stability Studies of Two Parallel B-Helix Proteins: Pectate Lyases C and E Renhao Li. Frances Jurnak and Clare Woodward

Crystal Structure of the Hemopexin-like C-terminal Domain of Gelatinase A

Andrew W. Libson, Apostolos G. Gittis, Ivan E. Collier, Barry L. Marmer, Gregory I. Goldberg and Eaton E. Lattman

Interactions of Cl-921, An Anilinoacridine Threading Agent with Nucleic Acids Ching-Yi Lien and David E. Graves

Characterization on Structure and Thermodynamic Stability of the C-terminal RBD2 of Human U1A Protein

Jirong Lu and Kathleen Hall

Site-Directed Mutagenesis Studies of the Stability of the Sac7d DNA-binding Protein from the Hyperthermophile Sulfolobus acidocaldarius

Branford S. McCrary, Stephen P. Edmondson and John W. Shriver

Thermodynamic Studies of Lac Repressor-Operator Interactions

Sonya E. Melcher, Diane E. Frank, Mark M. Levandoski, Oleg V. Tsodikov, Ruth S. Spolar and M. Thomas Record, Jr.

Core packing defects in an engineered Cro monomer corrected by combinatorial mutagenesis.

A.K.M.M. Mollah and Michael C. Mossing

Fluorescence Studies of the Ligand Induce Conformational Changes in the *E. coli* Hsp70, DnaK

Diana Montgomery & Richard Morimoto

Determination of Rate Constants using HPLC Affinity Chromatography

Peter D. Munro and Gary K. Ackers

Proton Linkage In Isothermal Titration Calorimetry

Kenneth P. Murphy and Brian M. Baker

Valence screening of water in protein crystals reveals potential Na+binding sites

Murad Nayal and Enrico DiCera

Purification and Characterization of Intact and Truncated Forms of the Biotin Accepting Subunit of the Escherichia coli Acetyl-CoA Carboxylase

E. Nenortas and Dorothy Beckett

Non-random Structure in Reduced and Unfolded Bovine Pancreatic Trypsin Inhibitor Hong Pan, Elisar Barbar, George Barany and Clare Woodward

Crystallographic, Molecular Modeling and Biophysical Characterization of the Valine B67 (E11)-Threonine Variant of Hemoglobin

Igor Pechik, Xinhua Ji, Krzystof Fidelis, Michael Karavitis, John Moult, William Brinigar, Clara Fronticelli and Gary L. Gilliland

DNA Sequence-Specific Differences in the TBP-Promoter Interaction

Victoria Petri, Elizabeth Jamison and Michael Brenowitz

The trp Repressor of E. coli: Further Physical Investigations of Super-Repressor Mutants

Ross J. Reedstrom, Shyam Vangala, Kathleen S. Martin, Erik W. Wilker & Catherine A. Royer

Interaction of Netropsin and Distamycin with the Grooves of an Intramolecular DNA Triplex

Dionisios Rentzeperis & Luis A. Marky

Solvent Exchange of the Tyr 31 Hydroxyl Proton in Ovomucoid Third Domain: A Sensitive Measure of the Effect of Protein Structure on Hydroxyl Acid/Base Character

Andrew D. Robertson and Junmin Peng

Allosteric Modulation by Tertiary Structure in Mammalian Hemoglobins. Introduction of the functional characteristics of bovine hemoglobin into human hemoglobin by five amino acid substitutions.

Maria Teresa Sanna, Michael Karavitis, William S. Brinigar and Clara Fronticelli

The application of synchrotron x-ray radiation to the development of time resolved methods for the study of RNA folding

Bianca Sclavi, Sarah Woodson, Mike Sulliven, Mark Chance and Michael Brenowitz

Calcium-Induced Differences in the Hydrodynamic Behavior of Calmodulin Mutants Explored by Gel Permeation Chromatography

Brenda R. Sorensen and Madeline A. Shea

Excluded Volume Effects on the Melting of DNA Duplexes and Triplexes

C.H. Spink and J.B. Chaires

Temperature and pH Dependences of Hydrogen Exchange and Global Stability for Ovomucoid Third Domain

Liskin Swint-Kruse and Andrew D. Robertson

RNA Polymerase-Promoter Kinetics and Thermodynamics: Results of an Analytical Method

Oleg V. Tsodikov, Peter E. Schlax Jr., Kristi L. McQuade and M. Thomas Record, Jr.

Investigation of Protein-Protein interactions of trp super-repressor mutants Shyam S. Vangala, Kathleen S. Martin, Ross J. Reedstrom and Catherine A. Royer

Proline Isomerization and the High-Pressure Denaturation of Staphylococcal Nuclease

Gediminas A.J. Vidugiris, Dagmar M. Truckses, John L. Markley and Catherine A. Royer

Role of the Peptide Backbone in Stabilization of Proteins by the Counteracting Osmolyte, Trimethylamine-N-Oxide

Aijun Wang and D. W. Bolen

Effects Of A Naturally Occurring Compatible Osmolyte On The Internal Dynamics of Ribonuclease A

Aijun Wang, Andrew D. Robertson, D. W. Bolen

Physicochemical Characterization of RNA Stem Loops with Non-nucleotide Spacers
D. Jeremy Williams & Kathleen B. Hall

Construction And Characterization of the ATPase Activity of a Covalently Cross-Linked Rep-DNA Complex

Isaac Wong and Timothy M. Lohman

Evidence for Inter-Domain Interaction in the Escherichia coli Repressor of Biotin Biosynthesis from Studies of an N-terminal Deletion Mutant

Yan Xu and Dorothy Beckett

Thermodynamic Analysis of Binding of the Escherichia coli Repressor of Biotin Biosynthesis to Small Ligands

Yan Xu and Dorothy Beckett

A Proton Binding Study of Thermodynamic Characteristics of the Denatured States of Staphylococcal Nuclease Proteins

Mingli Yang and Wayne Bolen